



CeNS/SFB1032 Workshop 2014 Walk and Talk at the Nanoscale

September 22 - 26, 2014 Venice International University (VIU), San Servolo, Italy



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Monday, September 22 (morning session)

Force Scaling in Stress Fibers

Timothée Vignaud, Laetitia Kurzawa, Ben Fogelson, Fabrice Senger, Jonathan Arnaud, Jean-Louis Martiel, Alex Mogilner, Laurent Blanchoin, <u>Manuel Théry</u>

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Cells have the remarkable ability to sense geometrical and physical cues from their environment and adapt their architecture accordingly. This process requires a tight regulation of the permanent remodeling of the acto-myosin network, that can both transmit and generate intra-cellular forces. Despite numerous works on the molecular composition of stress fibers, little is known about the mechanism determining the magnitude of force production in these structures. Here we studied the scaling of contractile force magnitude in stress fibers and investigated the role of actin network dynamics and architecture in this process. We used micropatterned substrates to control the length and spatial organization of stress fibers in adherent cells and measured the traction forces they produced on deformable substrates. Thereby, we demonstrated that forces scaling exhibit a biphasic behavior. Force magnitude first increased with the length of the stress fibers and then dropped above a critical length. Strikingly, very long cells appeared capable to produce only weak forces. Monitoring stress fiber relaxation upon laser nano-surgery, we showed that stress fibers were connected to the surrounding actin meshwork all along their length. A theoretical model accounting for the biphasic behavior of force scaling established friction between actin stress fibers and their surrounding cytoskeleton as being a key parameter in the regulation of force production by the cells.

Approaching the limit: Multiplexed Super-Resolution Microscopy with DNA-PAINT and Exchange-PAINT

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Super-resolution fluorescence microscopy is a powerful tool for biological research, but obtaining multiplexed images for a large number of distinct target species remains challenging. Here we use the transient binding of short fluorescently labeled oligonucleotides (DNA-PAINT, a variation of point accumulation for imaging in nanoscale topography) for simple and easy-toimplement multiplexed super-resolution imaging that achieves sub-10-nm spatial resolution *in vitro* on synthetic DNA structures. We report a multiplexing approach (Exchange-PAINT) that allows sequential imaging of multiple targets using only a single dye and a single laser source. We experimentally demonstrate ten-color super-resolution imaging in vitro on synthetic DNA structures as well as four-color two-dimensional imaging and three-color 3D imaging of proteins in fixed cells.



Can thermal traps drive Darwinian evolution?

Christof B. Mast¹, Matthias Morasch¹, Severin Schink², Ulrich Gerland² and Dieter Braun¹

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Long and complex biopolymers like RNA, DNA and proteins are essential for life on earth. Their emergence and replication are the prerequisites for early molecular Darwinian evolution. However, the origin of the first RNAzymes remains unclear because hydrolytic dissociation limits the maximal length of prebiotic polymers in aqueous, even millimolar concentrated nucleotide solutions to ~20 bases. While effective polymerization of RNA is possible in dry conditions [1], we demonstrate how a simple thermal gradient across an elongated, water-filled pore enhances a hybridization-based, worst-case polymerization process [2]: The thermal gradient drives a fluid convection and an orthogonal movement of biomolecules, also known as thermophoresis. Both effects lead to a massive accumulation of biomolecules at the pore bottom. This thermal molecule trap concentrates longer polymers more effectively than short monomers while polymerization works the better, the higher the total monomer concentration is. In combination, both processes are mutually self-enhancing and lead to an escalation of polymerization. We developed a theory for trapped polymerization and experimentally validated it using the hybridization based polymerization of dsDNA in a laser driven thermal trap. Extrapolation of the theory toward the RNA-world shows that a pore height of 5 cm and a temperature difference of 10 K are sufficient to form RNA polymers longer than the shortest RNA based replicator. In the experiments, we found that the escalation of polymerization also leads to a sequence selective phase transition of diluted oligomers into a gel-like DNA complex. Only polymerizing monomers with matching sequence are gelated while non-polymerizing monomers remain in the diluted state. The millimeter-sized, highly concentrated complexes remain stable

Graphene Plasmons: Properties and Applications Phaedon Avouris

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I will begin with a short introduction to the electronic structure and the single particle excitations of graphene. The focus will then be on the collective excitations (plasmons) of this material. Next will be discussed the properties of graphene plasmons, comparing them with those of the surface plasmons of conventional metals. I will mostly concentrate on localized graphene plasmons in lithographically patterned nano- and micro-structures (quantum dots and ribbons). I will discuss their optical even without active trapping and could have enhanced prebiotic, RNAzyme-catalyzed reactions. At the same time, the oligomer gel could have protected its selected constituents against diffusion without the help of lipids or fatty acids. We show that thermal traps can also drive the exponential replication of genetic information which is essential for Darwinian evolution ^[3]. The convective fluid flow thermally circulates oligomers and replicators, while replication products are accumulated and protected against outward diffusion into the diluted reservoir. In a proxy replication reaction, DNA replicating polymerase is able to double the amount of a 143mer product each 50 s, while the time constant for accumulation is 92 s. Thermal traps could therefore represent a possible non-equilibrium environment for the formation, selection and replication of the first biopolymers.

[1] Dry polymerization of 3',5'-cyclic GMP to long strands of RNA. Matthias Morasch, Christof Mast, Johannes Langer, Pierre Schilcher and Dieter Braun (2014) ChemBioChem 15,6:879-883

[2] Escalation of polymerization in a thermal gradient. Mast CB, Schink S, Gerland U, Braun D (2013) Proceedings of the National Academy of Sciences 110:8030–8035.

[3] Thermal Trap for DNA Replication. Mast CB, Braun D (2010) Physical Review Letters 104:188102

behavior in the infrared and THz regions of the spectrum, size effects, doping effects, effects of external magnetic fields, and the hybridization of the graphene plasmons with substrate and adsorbed overlayer optical phonons and their damping mechanisms. Applications of graphene plasmons in passive THz optical elements, the enhancement of photocurrents in graphene infrared photodetectors, and the enhancement of infrared absorption spectra of molecules will be discussed.

Chemical Strategies for Delivery of RNAi Drugs

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Chemically engineered synthetic small interfering RNA siRNAs act as therapeutic agents through the RNA interference (RNAi) pathway and are specific and potent inhibitors of gene expression. These agents may be designed to target disease pathways previously considered "undruggable". Numerous proof-of-concept studies both in animal models of human disease and in clinical trials demonstrate the broad potential and therapeutic value of RNAi therapeutics. The major challenge for the successful development of systemically delivered RNAi therapeutics had been the efficient delivery into organs or tissues and cells of interest to elicit RNAi mediated knockdown of faulty genes and effective translation of these approaches into clinic. Efficient delivery approaches have been developed over the years and clinical trials are advancing with RNAi therapeutic candidates formulated in lipid nanoparticles (LNPs) for intravenous administration.

We also have developed multiple conjugation strategies with the goal of robust *in vivo* delivery of siRNAs for therapeutic use. Covalent conjugation of small molecules to siRNA may avoid side effects resulting from the use of non-viral vectors, particles, or excipient-based delivery systems. Chemical modifications, mode of administration, and the nature of the conjugated ligand play significant roles in systemic delivery of siRNAs. In our early work with cholesterol and other lipophile-conjugated siRNAs, we demonstrated that covalent conjugation of small molecules can enhance bioavailability of siRNAs to multiple tissues including liver. Uptake of siRNA-lipophile conjugates is facilitated by lipoprotein particles and associated receptors and by other transmembrane proteins that are ubiquitously expressed. Systemic and local applications of cholesterol-siRNA conjugates for various therapeutic targets have also been broadly documented.

More recently, systemic delivery of therapeutic siRNAs to liver hepatocytes by subcutaneous administration has been achieved by conjugating chemically modified siRNAs with multivalent Nacetylgalactosamine (GalNAc) residues that are recognized by the asialoglycoprotein receptor (ASGPR). ASGPR is a C-type lectin receptor expressed on the cell surfaces of mammalian hepatocytes at 0.5 to 1 million copies per cell and is highly conserved across species. ASGPR recognizes an exposed terminal galactose (Gal) and enables clearing of serum glycoproteins via clathrin-mediated endocytosis. Recognition by the ASGPR requires multi-valency with appropriate spatial orientation of the Gal or the high affinity sugar analog N-acetylgalactosamine (GalNAc). Binding affinities of these ligands vary from mM to low nM and depend on the number of sugar residues present. siRNA-GalNAc conjugates efficiently target and silence disease-causing genes produced in liver hepatocytes. Using this conjugation platform, Alnylam is advancing several RNAi agents specific for liver targets through pre-clinical and clinical development to address genetically defined diseases with highly unmet medical need. Our progress with the chemistry of siRNA-GalNAc conjugates and applications in several therapeutic areas will be presented.

Precision Positioning and Sensing for Nano-Manipulation Applications Khaled Karrai

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Precision positioning and precision position sensing in space constrained is particularly challenging. It becomes even more challenging to address applications demanding specific manipulation in environmental constraints, such as cryogenic temperatures, or ultrahigh vacuum. Here I review and discuss by way of example few of such applications. One of them concerns the fabrication of quantum dot based quantum devices. Pioneering novel techniques in semiconductor device fabrication have recently greatly benefitted from the possibility optical lithography to make novel quantum optical devices out of randomly distributed quantum optical light emitter ^[1-3]. For instance by exposing an appropriate photoresist while optically localizing the position of light emitting single quantum dots allowed high-yield fabrication of novel quantum optical devices. Here, as an example, once the single quantum dot light emitters are spatially individually localized and their emission color identified, appropriate optical devices are subsequently lithographically defined to imbed each quantum emitter individually to insure the device full functionality. This so called deterministic lithography approach allows coping with the typical randomness of the spatial and spectral distribution of single semiconductor quantum dots demonstrating a rare case of confluence between bottom-up and top-down approaches in nano-device technology. The emission wavelength and intensity of single semiconductor quantum dots happens to be strongly temperature dependent and it is crucial that such deterministic optical lithography takes place at cryogenic temperatures ranging between 4K and 100K. In this work we show how the development of piezo displacement stages and accurate position readout allowed for building a cryogenic optical lithographer.

[1] A. Dousse et al., Phys.Rev.Lett.101, 267404 (2008)

[2] A. Dousse et al., App.Phys.Lett. 94, 121102 (2009)

[3] A. K. Novak et. Al. Nature Communication (Feb. 2014)

Plasmonic metal oxide nanocrystals and their near infrared electrochromism Delia J. Milliron

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Degenerately doped metal oxide semiconductors exhibit plasmonic resonance at wavelengths tunable by varying their composition. We have recently developed robust chemical strategies for incorporating high levels of dopants during colloidal nanocrystal synthesis and thereby fabricated a wide range of dispersible, processible nanocrystals with infrared plasmon absorption features.^[1] For example, aluminum-doped zinc oxide nanocrystals exhibit broad absorption peaks in the mid-infrared that change intensity and position systematically with doping level, while plasmon absorption peaks of tin-doped indium oxide (ITO) nanocrystals can be tuned across the near infrared. These plasmonic metal oxide nanocrystals are chemically robust, which has enabled us to study their optical response to electrochemical charging and discharging. Over a few volts of applied bias, the carrier concentration and plasma frequency in a conductive network of ITO nanocrystals were reversibly modulated by nearly 3- and 2-fold, respectively.^[2] The result is a strong and selective modulation of near infrared transmittance in visibly transparent films. This effect shows promise for energy-saving smart windows, a potential application which is even

more compelling when considering the electrochromic properties of inorganic composites containing these nanocrystals. ITO-in-amorphous niobium oxide composites exhibit sequential switching of near infrared and visible transmittance whose dynamic range is enhanced 5-fold by synergistic reorganization at the ITO-glass interface.^[3] Our recent prototype "smart glass" devices demonstrate the opportunity to dynamically control solar heat gain and daylighting in buildings in response to changing environmental conditions. I will conclude by highlighting outstanding fundamental questions about plasmons in metal oxide nanocrystals^[4] and their broad range of possible applications.

[1] R. Buonsati and D. J. Milliron, Chem. Mater., 2013, 25, 1305

[2] G. Garcia et al., Nano Lett., 2011, 11, 4415

[3] A. Llordes et al., Nature, 2013, 500, 323

[4] S. D. Lounis et al., J. Phys. Chem. Lett., 2014, 5, 1564

On the use and abuse of THERMODYNAMIC entropy Peter Hänggi

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Let us elaborate on the notion of *thermodynamic* entropy S (Clausius 1865) and its consequences. Gibbs put forward two notions of entropy for isolated systems that I commonly will refer to as `volume entropy' (involving the integrated density of states in modern language) and as the `surface entropy', being proportional to the density of states, commonly also known (incorrectly) as the Boltzmann entropy. The absolute temperature, $T^{-1}=\rhoartial S/\rhoartial U$, is then related to thermodynamic entropy; -- but which one to use? -- The consistency for thermodynamics, i.e. the validity for the celebrated 0-th, 1-st and 2-nd thermodynamic Law singles out the Gibbs-entropy^[11].

I shall address shortcomings that relate to the thermodynamics of small systems when sticking to the (Boltzmann)-surface entropy^[2,3]. Most of all, the uncritical use of Boltzmann entropy for microcanonical systems may formally yield negative values of absolute temperatures. This is not only physically incorrect for the concept of an *absolute* temperature, but also would violate thermodynamic stability if the system is brought into (weak) contact with an omnipresent sort of environment of radiation source or otherwise with no upper bound in energy. This criticism applies to the concept of absolute negative (spin) temperature and, as well, to the interpretation of recent experiments with isolated ultra-cold atomic gases^[4].

Next, we address canonical entropy when describing quantum systems that interact {\it strongly} with an environment. Then, the canonical specific heat can assume negative values away from absolute zero temperature^(5,6). Likewise, the thermodynamic entropy for a strongly coupled system, assuming a form which mimics a conditional entropy (but not quite) can be negative away from absolute \$T=0\$.

[1] S. Hilbert, P. Hänggi, and J. Dunkel, Thermodynamic Laws in Isolated Systems, arXiv:1408.5382 (2014).

[2] M. Campisi, Microcanonical phase transitions in small systems, arXiv 0709:1082; ibid, On the mechanical foundations of thermodynamics: The generalized Helmholtz theorem. Studies in History and Philosophy of Modern Physics 36, 275 (2005)

[3] J. Dunkel and S. Hilbert, Phase transitions in small systems: microcanonical vs. canonical ensembles, Physica A 370, 390 (2006).

[4] S. Braun, et al., Negative absolute temperatures for motional degrees of freedom, Science 339, 52 (2013); J. Dunkel and S. Hilbert, Consistent Thermostatics forbids Negative Absolute Temperatures, Nature Physics, 10: 67-72 (2014).

[5] P. Hänggi, G.L. Ingold, and P. Talkner, Finite quantum dissipation: the challenge of obtaining specific heat, New J. Phys. 10, 115008 (2008); ibid, Specific heat anomalies of open quantum systems, Phys. Rev. E 79, 061105 (2009).

[6] M. Campisi, P. Talkner, and P. Hänggi, Thermodynamics and fluctuation theorems for a strongly coupled open quantum system: an exactly solvable case, J. Phys. A: Math. Theor. (Fast Track) 42, 392002 (2009).

See also the two links:

(1) "(Quantum)-FluctuationTheorems" http://www.physik.uni-augsburg.de/theo1/hanggi/Fluctuation.html

(2) "What is Temperature" http://www.physik.uni-augsburg.de/theo1/hanggi/Temperature.html

Molecular computing meets synthetic biology^[1] Yaakov Benenson, ETH Zurich, Switzerland

One of the motivations behind computing with molecules is to "computerize" living systems, for example to prevent disease or control artificial tissues. Biology, however, is already very good at computing - the human brain being one example. Even on a single cell level information is constantly being processed, and the development of a functional organism from a single fertilized cells is controlled by an ingenious if only partially understood program encoded in DNA. Does this mean that the efforts to "write" new molecular programs are redundant? Not at all - natural programs have taken three billion years to evolve and, despite their beauty, are very difficult to alter in any way.

In my view the optimal approach is to balance the engineering principles inspired by computer science and engineering such as universal models, reprogrammability, modularity, etc., with the harsh reality of cell and organismal biology. The simple fact is that we do not know yet, even at the theory level, whether it is possible to perform reliable information processing in actual living cells as opposed to idealized "well-mixed reactors". Despite these limitations, the field of molecular computing in cells, or biological computing, has made significant steps forward with new design principles, new architectures, and new exciting experimental results. These developments also inform basic biological research.

[1] Unconventional Computation and Natural Computation. 13th International Conference. Proceedings: LNCS 8553. Publisher: Springer, Cham, Switzerland

Studying cellular structure and function with 3D structured illumination microscopy (3D-SIM) and fluorescent nanobodies

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Fluorescence light microscopy allows multicolor visualization of cellular components with high specificity, but its utility has until recently been constrained by the intrinsic limit of spatial resolution. We applied three-dimensional structured illumination microscopy (3D-SIM) to circumvent this limit and to study the mammalian nucleus (*Science, 320, 1332-6*). By simultaneously imaging chromatin, nuclear lamina, and the nuclear pore complex (NPC), we observed several features that escape detection by conventional microscopy. We are now studying the topological organization of nuclear functions at super-resolution. Multicolor 3D-SIM opens new possibilities to analyze subcellular structures beyond the diffraction limit of the emitted light. We have recently generated fluorescent, antigen-binding proteins, termed chromobodies, combining epitope-recognizing fragments with fluorescent proteins (*Nature Methods, 3, 887-*9). Unlike conventional antibodies these chromobodies can be expressed in living cells and used to target or trace epitopes in subcellular compartments. As chromobodies are soluble and active in the intracellular environment, they provide an optical readout for novel high content analyses and enable functional studies in vivo (*Nature Struct. Mol. Biol., 17, 133-139*). These antigen-binding fragments can also be produced in *E. coli*, chemically functionalized and used for super-resolution microscopy (*Science, 331, 1616-20*).

Quantum emulation with microcavity polaritons

Alberto Amo

CNRS-Laboratoire de Photonique et Nanostructures, Marcoussis, France

Semiconductor microcavities are an excellent platform to study non-linear optical phenomena and the quantum fluid properties of Bose-Einstein condensates. The eigenstates of this system are polaritons, half-light/half matter quasiparticles arising from the strong coupling between excitons and photons confined in an optical cavity of micrometric size. Thanks to their very light mass (10⁻⁴ times the electron mass) these photonic bosons can form Bose-Einstein condensates at temperatures much higher than in atomic gases (10K for polaritons vs 100nK for atoms). By engineering the shape of the microcavities we can use polaritons to emulate various Hamiltonians in a photonic system directly accessible using standard optical techniques. Taking advantage of these properties we have experimentally studied Josephson oscillations and photonic self-trapping in a microcavity, and fabricated a honeycomb lattice in which photons behave like electrons do in graphene.



Polariton honeycomb lattice



Single-molecule studies of genome processing Nynke Dekker

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Over the past few decades, there has been steady progress in both our ability to produce biological material and in our ability to manipute matter at small length scales. These two developments merge in a fascinating area of confluence called singlemolecule biophysics in which an understanding of biological matter from physical principles becomes possible. I will

illustrate how several newly-developed single-molecule force and torque spectroscopy techniques allow us to shed light on genomic processes such as transcription, replication, and DNA compaction. Lastly, as the true environment of biological molecules is the living cell, I will demonstrate our ability to track replication inside bacterial cells, and discuss implications for the future.



TUESDAY, SEPTEMBER 23 (AFTERNOON SESSION)

Targeted nanocomplex formulations for gene and siRNA therapy

Stephen Hart

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Synthetic, non-viral vectors such as cationic liposomal and polymeric formulations offer advantages over viral vectors for in vivo gene therapy in that they are less immunogenic, have fewer packaging constraints and are safe. However their transfection efficiencies in vivo are usually insufficient. Multifunctional synthetic formulations are being developed that emulate the properties and functionalities of viral capsids and higher transfection efficiencies. We are developing a formulation that we term a Receptor Targeted Nanocomplex (RTN), which is a mixture of cationic, receptor-targeting peptides and liposomes with nucleic acids, plasmid DNA (pDNA) or short interfering RNA (siRNA). The peptide mediates DNA packaging and receptor targeting while the lipid fuses with the endosomal membrane leading to improved cytoplasmic release of the DNA or siRNA. RTNs display a synergistic enhancement of transfection due to the combined functionality of these components, both for siRNA and plasmid delivery.

Diseases affecting different tissues are likely to need specifically designed formulations optimised for the target tissue and the route of delivery. RTN formulations have been developed with anionic or cationic charges, and with stealth properties resulting from PEGylation. The modularity of the RTN design enables development of formulations optimised for specific therapeutic strategies including lung disease in cystic fibrosis with delivery by nebulisation, neuroblastoma cancers by systemic administration, and brain delivery for neurodegenerative diseases by direct injection.

Phase coexistence and charge traps in organic semiconductors – enlighten the disorder at the nanoscale –

Christian Westermeier^{1,2}, Adrian Cernescu³, Sergiu Amarie³, Clemens Liewald¹, Fritz Keilmann¹, Bert Nickel^{1,2}

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The control of nanocrystallinity and disorder in organic thin films such as pentacene is crucial for the improvement of electronic applications. The distribution of crystalline phases and trap states at the emerging interfaces is, however, difficult to access experimentally. We have successfully developed a new imaging technique for trap densities in pentacene thin films by using essentially a pulsed laser scanning microscope to scan across the channel of an organic field-effect transistor during operation.⁽¹¹⁾ Here, filling of the trap states can be adjusted via the gate volt-



subsequently age. the trapped charges are locally released by exciton-mediated trap clearing upon illumination and the released charge is detected as light-induced change of the transistor current. As photoresponse this is sensitive to the amount of trapped charges within the

The grain boundaries between two coexisting phases in organic semiconductor pentacene are expected to obstruct charge transport in its thin-film devices. We use infrared-spectroscopic nano-imaging to show an interlocking morphology, which is uncorrelated with its grain structures. illuminated area, the recorded map mirrors the distribution of traps throughout the pentacene film and hotspots can be identified. While the laser focus of our trap imaging technique is diffraction limited, we also utilize s-SNOM imaging that allows for local excitation of an organic film with a resolution down to 20 nm due to near-field enhancement via a metalized tip. This resolution, being independent of the wavelength, also holds for midinfrared light that excites molecular vibrations. We benefit from the sharp IR resonances of highly-ordered organic systems, which turn out to be sensitive even to the crystalline packing of molecules. Thus, IR-SNOM allows to image different crystalline phases of the same material in thin films for organic electronics. For the case of pentacene, we find that grains of a few microns size, which appear to be in the so called thin film phase, are indeed subject to massive nucleation of bulk phase pentacene.^[2] This may explain contradicting reports on the correlation of grain size and carrier mobility in pentacene films.

[1] C. Westermeier, M. Fiebig, B. Nickel, Mapping of Trap Densities and Hotspots in Pentacene Thin-Film Transistors by Frequency-Resolved Scanning Photoresponse Microscopy, Adv. Mater. 25(40): 5719–5724 (2013).

[2] C. Westermeier, A. Cernescu, S. Amarie, C. Liewald, F. Keilmann, B. Nickel, Sub-micron phase coexistence in small-molecule organic thin films revealed by infrared nano-imaging, Nat. Commun. 5: 4101 (2014).

Cooling and Amplification of a Vacuum-Trapped Nanoparticle

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We optically trap a single nanoparticle in high vacuum and cool its three spatial degrees of freedom by means of active parametric feedback. The small size and mass of the nanoparticle yield high resonance frequencies and high Q-factors along with low recoil heating, which are essential conditions for ground state cooling and for low decoherence. The vacuum-trapped nanoparticle forms an ideal model system for studying non-equilibrium processes, nonlinear interactions, and ultrasmall forces.

Figure 1: (top) Photograph of light scattered from a trapped 85 nm fused silica particle (arrow). The object to the right is the outline of the objective lens. (bottom) Time trace of the particles x coordinate (transverse to optical axis) at 2mbar pressure. Trapping times of several days have been achieved.



Wednesday, September 24 (morning session)

0.7-Anomaly in Quantum Point Contacts: Correlations in 1D

Florian Bauer,^{1, 2} Jan Heyder,^{1, 2} David Borowsky,¹ Enrico Schubert,¹ Daniela Taubert,¹ Dieter Schuh,³ Benedikt Bruognolo,^{1, 2} Werner Wegscheider,⁴ Jan von Delft,^{1, 2} and <u>Stefan Ludwig</u>¹

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3 Institut für Angewandte Physik, Universität Regensburg, Germany

4 Laboratory for Solid State Physics, ETH Zürich, Switzerland

Quantum point contacts (QPCs), the ultimate building blocks of quantum electronic circuits, are 1D constrictions in a 2D electron system (2DES). When a QPC is pinched off, its conductance famously decreases in integer steps of the conductance quantum, $G_Q = 2e^2/h$. The source of an unexpected feature with an intriguing dependence on temperature, magnetic field and source-drain voltage has been debated for the last two decades ^[1]. Often entailing a kink of the pinch-off curve near $0.7G_0$ it is called the 0.7-anomaly. In this talk I will explain how the 0.7-anomaly naturally arises from strong correlations fostered by the enhanced density of states in 1D at low energies. The correlations cause an anomalous increase of the spin susceptibility and the back-scattering rate. Our microscopic model is built on a combination of systematic measurements of a highly tunable QPC and detailed numeric calculations ^[2]. Compared to previous models we utilize no assumptions, such as a local moment or a broken time-reversal symmetry, in addition to experimental evidences.

[1] A. Micolich, J. Phys. Cond. Matt. 23, 443201 (2011)

[2] F. Bauer, et al., Nature 501, 73 (2013)



Scanning electron micrograph of our QPC sample (top) and animated potential view (bottom). A 2D electron system (2DES) resides 85nm beneath the surface of a GaAs/AlGaAs heterostructure. Two central gold gates (c), four side gates (s) and an electrically isolated global top gate (t) are used to shape by the electric field effect a 1D constriction in the 2DES. Electrons move through the QPC between source to drain contacts (dark blue areas) while they are repelled by high potential regions (yellow, red). [animation produced by Christoph Hohmann.]

Electron Interactions in Quantum Point Contacts

Hermann Sellier

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Quantum point contacts (QPCs) are well known for their waveguide properties, but they also show a rich physics of electron interactions. Applying a negative voltage on split-gates patterned on a two-dimensional electron gas (2DEG) creates a quasi onedimensional electron channel connecting two large reservoirs ^[1]. Each wave-guide mode carrying one conductance quantum 2e²/h, the conductance curve versus gate voltage shows a series of quantized plateaus which are well reproduced by a simple saddle potential model [2]. However, a shoulder-like feature is commonly observed at a conductance around 0.7×2e²/h which cannot be explained by single-particle theory [3]. With lowering temperature, this "0.7 anomaly" shades off and a "zero-bias anomaly" emerges [4]. Although the link between these structures remains an open question, both are thought to arise from strong Coulomb interactions in the small 2DEG region forming the QPC where the electron density is low. Different theoretical models have been proposed to explain these anomalies, but no consensus could be reached so far on their interpretation^[5].

After a brief review of experimental observations and theoretical models, I will present recent experiments 16,71 that indicate the presence of self-consistently localized charges in QPCs. In particular, I will focus on our experimental approach ^[7] using a Scanning Gate Microscope (SGM) [8-10] to change in-situ the shape of the QPC potential. The SGM tip is used as a movable gate in a more flexible and less invasive way than what could be achieved with several fixed surface gates. Approaching the tip towards the QPC produces an oscillatory splitting of the zerobias anomaly, correlated with simultaneous appearances of the 0.7 anomaly, thereby revealing that both features share a common origin. These repetitive changes are interpreted in terms of a many-body localized state, induced by the strong Coulomb repulsion in this low density region, and forming a small onedimensional Wigner crystal [11-13]. The number of charges in this crystal is controlled by the tip position and produces different Kondo screenings from the leads depending on charge parity, with a conductance peak either at zero, or at finite bias ^[14,15].

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Surface plasmons and phonon polaritons in atomically thin van der Waals crystals Dimitri N. Basov

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Layered van der Waals (vdW) crystals consist of individual atomic planes weakly coupled by vdW interaction, similar to graphene monolayers in bulk graphite. These materials can harbor superconductivity and ferromagnetism with high transition temperatures, emit light and exhibit topologically protected surface states. An ambitious practical goal is to exploit atomic planes of vdW crystals as building blocks of more complex artificially stacked structures where each such block will deliver layer-specific attributes for the purpose of their combined functionality. Infrared (IR) nano-spectroscopy and nano-imaging experiments on hexagonal boron nitride (hBN) have uncovered rich optical effects associated with phonon polaritons in this prototypical van der Waals crystal. We launched, detected and imaged the polaritonic waves in real space and altered their wavelength by varying the number of crystal layers in our specimens [*Dai et al. Science, 343, 1125, (2014)*]. Unlike surface plasmons in graphene that we have imaged using a similar nano-IR toolset [*Fei et al. Nature 487, 82 (2012)*], highly confined phonon polaritons are immune to electronic losses and therefore can travel over distances exceeding 10-s of microns. I will also discuss an ability to control plasmonic response of graphene at femto second time scales that we have demonstrated using a unique pumpprobe nano-IR apparatus [*Wagner et al. Nano Letters 14, 894 (2014*)].

Nucleic Acid Delivery – From Academic Discovery to Drug Development Christian Plank

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Nucleic acids carry the building plans of living systems and fulfil numerous further functions in living cells. In theory, any cellular function may be influenced in a purposeful manner if an appropriate nucleic acid can be shuttled into a target cell in a precise enough manner. In nucleic acid therapy, nucleic acids are used to complement or repair damaged genes or to interfere with endogenous gene expression. For this purpose, nucleic acids are usually formulated as multifunctional nanoparticles which are designed to overcome numerous barriers on the way from an administration site to the target cells. We have developed various methods for using nucleic acids in medical applications. More recently, we have focused on "gene therapy without genes", that is on using messenger RNA instead of genes for making patient cells produce their own therapeutic protein. This program is intended to enter first in man application in 2015. In this presentation, challenges encountered when developing nucleic acid drugs from bench to bedside will be discussed.

Complement Sensing at Nanoscale

S. Moein Moghimi

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The complement system is a complex network of plasma and membrane-associated proteins and represents one of the major effector mechanisms of the innate immune system. The function of complement in innate host defence is accomplished through highly efficient and tightly orchestrated opsonisation, lytic and inflammatory processes. Synthetic nanoparticles and particulate drug carriers (e.g., liposomes, polymeric nanoparticles and nanocapsules) by virtue of their size, shape and surface characteristics (e.g., display of architecture with repetitive epitopes such as those arising from surface projected polymers or polyanionic/polycationic clusters) may resemble many microorganisms. These 'pathogen-mimicking' properties make nanoparticles prone to interception by the complement system and a number of consequences ensue from complement activation. These comprise both beneficial and adverse reactions, depending on the extent and severity of complement activation as well as microenvironmental factors. These concepts will be discussed in relation to therapeutic applications of nanoparticles and anti-cancer nanomedicines.

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Nanoscale imaging and single-molecule detection at ultra-high concentrations using photonic antenna devices

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The quest for optical imaging of biological processes at the nanoscale has driven in recent years a swift development of a large number of microscopy techniques based on far-field optics. These, so called, super-resolution methods are providing new capabilities for probing biology at the nanoscale by fluorescence. While these techniques conveniently use lens-based microscopy, the attainable resolution and/or localization precision severely depend on the sample fluorescence properties. True nanoscale optical resolution free from these constrains can alternatively be obtained by interacting with fluorophores in the near-field. Indeed, near-field scanning optical microscopy (NSOM) using subwavelength aperture probes is one of the earliest approaches sought to achieve nanometric optical resolution¹. More recently, photonic antennas have emerged as excellent alternative candidates to further improve the resolution of NSOM by amplifying electromagnetic fields into regions of space much smaller than the wavelength of light. In this contribution, I will describe our efforts towards the fabrication of different nanoantenna probe configurations as well as 2D antenna arrays for applications in nano-imaging and spectroscopy of living cells²⁻⁵. I will show examples on how these devices can be exploited to detect individual molecules at micro-molar concentrations^{4,5}, as well as our efforts towards the study of cell surface receptors in living cells with unprecedented resolution and sensitivity.

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How yeast adapts to a strong genetic perturbation: one function at the time

Liedewij Laan and Andrew Murray

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Complex gene regulatory and protein networks allow yeast to grow and divide under a wide range of environments. These networks also need to be robust to genetic perturbations to buffer genotypic variation due to sexual reproduction and evolution. Nevertheless, on long evolutionary timescales mutations are essential to allow organisms to adapt. So, how robust are these networks to genetic perturbations, and what are the underlying mechanisms of robustness? We addressed these questions by deleting the nearly essential gene BEM1 from the polarity network in budding yeast. Bem1 brings together Cdc42, a small G protein that regulates actin polymerization, and its activator Cdc24, a guanine nucleotide exchange factor, and by doing so Bem1 is an essential part of one of the positive feedback loops that create a single site of cell polarization. Cells lacking Bem1 have severe polarization defects, resulting in many very large cells that eventually explode. As a consequence, $bem1\Delta$ populations proliferate roughly 10-fold slower than their wild-type counterparts.

To study adaptation, we evolved 11 $bem1\Delta$ lines by serial dilutions for one thousand generations. At the end of the evolution, bem1 Δ cells approached wild-type growth rate and cell size and polarization dynamics were indistinguishable from wild type. We sequenced the final time points and found mutations in three genes (BEM3, NRP1, and BEM2) that appeared in the same order in several independent lines. After reconstructing these mutations in a *bem1* Δ background and studying their phenotype, we suggest the following progressive mechanism of adaptation: The first mutation (inactivating BEM3) improves activation of Cdc42, the second mutation increases positive feedback, and the third mutation fine tunes activation of Cdc42. These experiments show that the polarization module is highly adaptable through a small number of mutations which are surprisingly efficient in recovering proper function, especially given the precision of the original network in its ability to produce one and only one site of polarization.

Strong spin-orbit coupling of spin-3/2 holes in gallium-arsenide semiconductor nanostructures Alex Hamilton

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The electrical current in a semiconductor can be carried by negatively charged electrons or positively charged holes. In undergraduate physics, we are often taught that holes in the valence band are just an absence of an electron. But they aren't. Valence band holes are spin-3/2 particles, and this gives them very different properties to spin-1/2 electrons. In recent years there has been growing theoretical interest in the possibility of using holes in semiconductor nanostructures for applications ranging from ultra-fast spin transistors through to quantum information and communication. This talk will describe where holes come from, why they are so different to electrons, and what one can do with holes that can't be done with electrons.

The differences between electron and holes are most striking when they are confined to low dimensional nanostructures. In quantum wires the interplay of spin-orbit interaction and electrostatic confinement leads to an extreme anisotropy of the Zeeman spin-splitting that is completely unlike electrons, and still not fully understood. I will present measurements mapping out the anisotropy of the hole Lande g-tensor, and show that it is possible to directly measure off diagonal component of the g-tensor.



Microfluidic droplets for quantitative biological studies

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I will present a set of tools that we have developed for manipulating microfluidic droplets. These tools simplify the fluidic operations, allowing the user to work with stationary drops while the conditions are varied in time. This gives access to spatial and temporal information on the contents of the drops, which can containt molecules (DNA, enzymes,...) or cells. Different applications of this platform have been developed in our lab. I will describe measurements of enzyme kinetics on an integrated chip, allowing us to obtain full enzyme activity characterisation rapidly and using minute amounts of reagents. I will also describe the detection of rare DNA through a digital PCR device based on our microfluidic tools. This device is now being industrialized by a spinout company (Stilla Technologies). Finally, I will describe a massively parallel cell culture platform that will allow the exploration of cecllular heterogeneity on a single chip. I will give examples of each of these devices and show some of the potential for doplet microfluidics for transforming the way biology is performed. I will also insist on the interest of the link between fundamental physical science and biotechnological applications.

Non-inertial lift and its application to label-free microfluidic cell separation and sorting <u>T. M. Geislinger</u>^{ab}, M. Stamp^a, B. Eggart^a, S. Braunmüller^a, L. Schmid^a, S. Chan^b, K. Moll^b, M. Wahlgren^b, A. Wixforth^a, and T.

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Reliable cell separation and sorting are important tasks in everyday's laboratory work and of increasing importance in various medical diagnoses. Widely used methods like fluorescence or magnetically activated cell sorting (FACS, MACS), however, require labelling of samples with adequate markers and/or the generation of external fields. Apart from the dimensions and the costs of such devices, any unwanted alterations of the cells by the markers potentially interfere with subsequent processes such as genetic analyses. Here, we present a simple and cheap microfluidic approach for continuous, passive and label-free cell sorting that relies on the exploitation of a hydrodynamic effect for separation: the non-inertial lift effect^[1]. The non-inertial lift effect is a repulsive cell-wall interaction of purely viscous origin that acts on non-spherical and deformable objects in laminar flow fields. Generally, the lateral drift becomes stronger for larger and more deformable objects. First, we examine the influence of flow rate and external fluid viscosity on the separation of different objects (red blood cells (RBC), blood platelets and latex microspheres) and compare our results with analytical theory^[2,3]. We then exploit our findings to separate RBCs from the smaller blood plate-

lets as well as blood platelets from the rigid, equally-sized latex microspheres^[2]. Based on our findings, we design a microfluidic device for non-inertial lift induced cell sorting (NILICS) that operates continuously and label-free, using size and deformability as intrinsic markers. We apply NILICS to sort circulating tumor cells out of RBC solutions by size with sorting efficiencies up to 100%^[4]. Furthermore, we use the much weaker dependency of the non-inertial lift on deformability to enrich RBCs infected with the malaria parasite *Plasmodium falciparum*^[5].

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Single-Molecule Fluorescence on DNA Origami

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In recent years, we have combined DNA nanotechnology with single-molecule fluorescence to create functional single-molecule devices such as nanoscopic rulers for superresolution microscopy and energy transfer switches^[1-4]. DNA origamis are also used to improve single-molecule detection by fluorescence enhancement with nanoantennas or by single-molecule placement in zeromode waveguides using nanoadapters^[5, 6]. Especially, fluorescence enhancement with self-assembled nanoparticles holds great potential for single-molecule detection at higher concentrations but also for diagnostic applications^[7, 8]. We will discuss recent advancement in fluorescence enhancement and how to disentangle the complex factors that influence the fluorescence of single molecules near metallic nanostructures^[9].

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Photo-Induced Charge Carrier Generation in Covalent Organic Frameworks

<u>Florian Auras</u>,^{a,b} Mona Calik,^{a,b} Laura Salonen,a Matthias Handloser,^{a,b} Mirjam Dogru,^{a,b} Dana Medina,^{a,b} Veronika Werner,^a Dirk Trauner,^{a,b} Achim Hartschuh,^{a,b} Paul Knochel,^a Thomas Bein^{a,b}

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Covalent organic frameworks (COFs) offer a strategy to position molecular semiconductors within a rigid network in a highly controlled and predictable way. The π -stacked columns of layered 2D COFs enable electronic interactions between the COF layers, thereby providing a path for exciton and charge carrier transport, while the accessible porous channels allow for infiltration of guest molecules. We have constructed ordered interdigitated heterojunctions by infiltrating thin films of a thienothiophene-containing COF with I6,61-phenyl C61 butyric acid methyl ester (PCBM) as electron acceptor (Figure 1). Photo-induced charge transfer from the COF to pore-located PCBM molecules was confirmed by strong photoluminescence quenching. Moreover, we used the COF:PCBM junctions as active layer in first COF-based photovoltaic devices.^[11] Another strategy for constructing opto-electroactive COFs, which we followed recently, is to combine two complementary semiconductors within the framework. If the energy levels and geometric arrangements are chosen adequately, the framework will consist of ordered columns of molecular donors and acceptors. We have applied this concept to construct a photovoltaic device based on a newly developed triphenylene-porphyrin COF. For the first time, we could demonstrate the generation of extractable charge carriers inside a COF upon illumination.

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Figure 1. a) The porous COF framework can be infiltrated with fullerene derivatives to create well-ordered interdigitated heterojunctions. b) Upon photoexcitation, electrons are injected into the fullerene phase and holes are transferred to the COF.

Imaging electron transport in semiconductor nanostructures on the scale of the electron wavelength

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Mesoscopic electron transport arises in semiconductor nanoand microstructures when the temperature is so low that several length scales, such as the elastic mean free path and the phasecoherence length of electrons become comparable to each other and to the system size. Since the field has matured over more than 30 years, a vast number of beautiful experiments and theories exist for understanding and describing the physics in these systems. Recently, an imaging technique has been developed that produces brilliant pictures of mesoscopic systems at the scale of the Fermi-wavelength of the electron. In this talk I will pick a few simple and basic examples from the available measurements to illustrate the imaging principle of this scanning gate technique and its physical meaning. We will see that some fundamental aspects of transport, such as conductance quantization, conductance fluctuations, or adiabatic edge channel transport in the quantum Hall regime can be rediscovered with spatial resolution and that there were unanticipated surprises that still leave open questions today.

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Effect of charge and size on molecule diffusion across basal lamina interfaces in vitro and in vivo	Sequence defined oligo(ethanamino)amides for folate receptor targeted pDNA or siRNA delivery: optimization and preliminary
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Harald Budde, Nicolás Coca López, Xian Shi.	Infrared hyperspectral manning of pano-composites
and Achim Hartschuh	Sergiu Amarie, Dominik Kalb, Fritz Keilmann, Clemens Liewald.
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Ultrastable Cellulosome-Adhesion Complex Tightens Under Load	Christian Westermeier
Constantin Schoeler, Klara H. Malinowska, Rafael C. Bernardi,	Magnetic (Torque) Tweezers Experiments to Probe
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POSTER ABSTRACTS - SESSION I (LI-Z)

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Effect of charge and size on molecule diffusion across basal lamina interfaces in vitro and in vivo <u>Fabienna Arends^{1,2,}</u> Sabine Sellner³, Markus Rehberg³ and Oliver Lieleg^{1,2}

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The permeability of the basal lamina, a biological hydrogel found at the basolateral side of the endothelium, is an important property for the design of new drug delivery systems. During inflammation, the endothelium becomes leaky and molecules can directly pass from the blood stream into the basal lamina. In such a situation, the basal lamina constitutes the main barrier against the efficient entrance of drug carriers from the blood stream into the connective tissue. Therefore, it is highly desirable to understand which parameters dictate diffusion processes across the blood-basal lamina interface. Here, we quantify the diffusion of dextran and peptides of different size and charge across a buffer-basal lamina interface using a microfluidics approach. As a model system for the basal lamina, we study an extracellular matrix gel (ECM) purified from the Engelbreth-Holm-Swarm sarcoma of mice. We measure the formation of a concentration gradient of solutes across the ECM phase and compare our findings with in vivo experiments obtained in living mice. From our data, we aim at deciphering the important parameters responsible for the permeability properties of the hydrogel. Our results can guide the design of new drug carriers which can efficiently diffuse from the blood stream into the connective tissue.

Covalent Organic Frameworks as Photoconductive Materials

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Covalent organic frameworks (COFs) have recently been developed as a new class of porous and crystalline materials with interesting and tunable physical and chemical properties.[1] Up to now, COFs can be achieved mainly by boronate ester formation, imine condensation or β -ketoenimine formation. Lately, two-dimensional COFs have attracted interest as potential candidates for the production of new photoactive materials.[2] The $\pi\text{-}interactions$ between the COF-layers allow for the formation of highly oriented films with host channels. These defined channels are accessible and can incorporate small quest molecules.[3] Here we present the two major approaches for the preparation of photoactive electron-donor-electron-acceptor COFs. The first route is based on the formation of highly ordered electron donor-acceptor interpenetrated systems, in which electron-donor COF films host electron-acceptor molecules such as C60 derivatives.[4] The second approach involves photoconductive COFs providing both electron-donating and electron-accepting subunits. Due to the π -stacked layers, these donor-acceptor-COFs

(D-A-COFs) exhibit highly ordered pillars of donor as well as acceptor entities that permit charge transfer within the COF and charge transport in the third dimension.[5]

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High-throughput force-fluorescence spectroscopy in zero-mode waveguides

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In the last years zero-mode waveguides (ZMW) emerged as an important tool to overcome the concentration limit of optical single-molecule detection. Via their subwavelength geometry in opaque metallic films they prohibit light propagation inside their cavity and hence are capable to confine excitation volumes of conventional optical microscopes drastically. This allows coping with high concentrations of fluorescently labeled ligands as necessary in single-enzyme experiments to monitor their biochemical activity (e.g. DNA real time sequencing). Complementary, mechanical experiments with single-molecule resolution and piconewton sensitivity provide control over a second but important parameter in molecular interactions: force. First combinations of these strong single-molecule properties, fluorescence and mechanics, provide enormous possibilities in mechanoenzymatic research. Our group addresses the employment of nanoapertures for AFM-based force spectroscopy to establish simultaneous single-molecule fluorescence-force spectroscopy at high fluorescent ligand concentrations. In a proof-of-principle experiment the analysis of a probable single binding event in a force-activated enzyme (Titin Kinase) has been shown in a ZMW. The methodology is further improved by the implementation of non-invasive tip localization routines to provide automated data acquisition at rates similar to those in standard force spectroscopy. Whereas light incident on the thin cantilever is transmitted with some losses, the fraction of light incident on its high-aspect ratio tip is strongly absorbed. Live superresolution methods applied on this absorption signal are able to securely and centrally navigate the cantilever into a ZMW with an accuracy of few nanometers. The application of high-throughput force-fluorescence spectroscopy in nanoapertures is promising for investigating the mechano-activation of kinases by high-yield recording of fluorescent ATP-binding or of minimum phosphorylation peptides under mechanical stress.

Traceless pH-sensitive coating of viral and non-viral vectors

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The successful implementation of oncolytic viruses or nucleic acids as medical agent depends on the delivery efficiency. Therefore, non-viral delivery systems have been developed to protect nucleic acids by compacting the negative charged nucleic acids into particles of virus-like dimensions. Alternatively, oncolytic virus vectors have emerged as novel natural tool to target and cause cytolysis of cancer cells [1]. Despite many advances on the vector development front, systemic delivery remains a major bottleneck. Nonspecific interactions with cells of the immune system as well as blood proteins and rapid clearance by liver and spleen have to be prevented. This can be overcome by the usage of shielding agents such as PEG (polyethylene glycol) and HPMA (N-(2-Hydroxypropyl)methacrylamide) [2]. However, irreversible surface shielding can lead to inactivation in case of oncolytic virus and endosomal entrapment in case of a polymeric carrier system. A promising method of improving the efficacy by overcoming these boundaries is the creation of vectors that are bioresponsive, so that transfer is enabled by cleavage of chemical bonds upon exposure to various physiological environments. Here, we present an approach where the delivery systems are reversible shielded with PEG or HPMA connected via a universal attachable tool called AzMMMan (azidomethyl-methylmaleic anhydride) [3], which can be attached to any carrier via primary amino groups. After the delivery systems enter an acidic environment (tumor tissue or endosome), a pH-labile bond is broken, releasing the functional virus or the vectors endosomolytic capability. We hypothesize that this approach proposed here will significantly enhance the transduction efficiency of systemically applied viral and non-viral delivery systems by avoiding unspecific interactions, hence prolonging circulation and therefore providing a substantial benefit to the clinical translation of these vectors as potent medical agent.

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Small Angle X-Ray Scattering for Structural Analysis of Biological Macromolecules Linda Bruetzel and Jan Lipfert

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Small angle X-ray scattering (SAXS) is a powerful technique to unravel the structure and interactions of biological macromolecules such as proteins and nucleic acids with a size of kDa up to GDa. In contrast to X-ray crystallography, the molecules are studied in solution allowing for time-resolved investigations and measurements under a broad range of solution conditions, including (near-) physiological environments. Thereby, only small sample volumes (10 - 30 µl) and moderate protein concentrations (~ 1 mg/ml) are required. SAXS experiments are routinely performed at state-of-the art 3rd generation synchrotron X-ray sources. As a complimentary approach we present an in-house setup for SAXS measurements, which we are currently developing in collaboration with the lab of Bert Nickel and Joachim Rädler. As a proof-of-concept, we are employing both double stranded DNA and proteins such as cytochrome c and lysozyme used previously as scattering standards.

Using SAXS we want to study the conformational changes of the large blood glycoprotein von Willebrand factor (vWF), which plays an important role in hemostasis and thrombosis. Recent studies could show that vWF undergoes regulating conformational changes at different hemodynamic conditions. SAXS measurements have the potential to provide further insight into pH-dependent structural dynamics of vWF. In addition, we describe how SAXS data can be used for the ab inito reconstruction of low-resolution (30-10 Å) 3-D models of biomolecules and their assemblies. In this context, we will perform SAXS measurements in order to characterize nucleic acids (RNA, DNA). In addition, gold nanocrystals, which can be attached site-specificly to the DNA will be utilized in order to obtain molecular distance distributions.

Surface acoustic wave induced charge carrier dynamics in single GaAs/AlGaAs radial heterostructure nanowires

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We investigate the influence of the dynamic electric fields induced by a surface acoustic wave (SAW) on the optical emission of single semiconductor nanowires (NW) via the use of microphotoluminescence (μ -PL) spectroscopy. The investigated nanowires consist of a GaAs core surrounded by an AlGaAs shell containing an additional 3nm wide radial GaAs quantum well (QW). For both, the core and the radial QW, we observe clear signatures of acousto-electric exciton dissociation and conveyance of electrons and holes. This method enables us to locally and contactlessly observe the electrical field induced carrier dynamics and make conclusions about the underlying mobilities of electrons and holes for both the core and the quantum well of the nanowire. Moreover, at high laser excitation powers and thus high charge carrier concentrations we observe an attenuation of this effect pointing to a screening effect of the piezoelectric fields caused by an increase of the charge carrier density.

Photophysics of hybrid organo-halide lead perovskite nanoparticles

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Hybrid organo-halide lead perovskites have raised great expectations in recent years due to their application as light harvesters in solar cells, resulting in promising energy-conversion efficiencies which are being improved at a remarkably fast pace. The simple and cost-effective construction of these devices appears as a viable option over current technologies which require the energy-intensive fabrication of highly crystalline materials. To understand the principles behind the performance of organohalide lead perovskites, it is fundamental to study their photophysical properties at the single crystal level. For this purpose, we have synthesized perovskite nanoparticles by means of a simple colloidal precipitation process. Homogeneous solutions of the precursor ions are slowly destabilized by the addition of poor solvents which are miscible with the continuous medium. This triggers the instantaneous formation of nano-sized crystals in a metastable colloidal state, which are immediately fixed to a substrate before further growth and destabilization takes place. The variation of concentrations and solvent ratios results in the formation of crystals of different sizes. This allows the photophysical study of crystals and assemblies of different sizes by means of single-particle photoluminescence spectroscopic techniques.

Dose response relation of triplebody mediated cell killing studied in single cell arrays

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Monoclonal antibody therapy is a promising strategy to treat cancer with the help of the patient's own immune system. Recent studies in the case of acute myeloid leukemia (AML) have shown that single chain triplebodies that recognize two epitopes on the cancer target cell provide more effective elimination. Here we study the potency of triplebody (SPM-2) to direct NK cells against cancer using single cell arrays of model target cells. Using fluorescence indicators and time-lapse microscopy the fraction of cells killed in the presence of NK cells is determined in an automated high-throughput modality. The approach allows to determine the killing fraction as a function of SPM-2 concentration as well as a function of effector-target (ET) cell number ratio. Our results demonstrate the potential of micro-structured cell arrays for quantitative studies on the interplay between the immune system and cancer cells and its antibody mediated manipulation.

Fourier Space Imaging of Raman Scattering Intensities in Graphene

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Raman spectroscopy has proven to be a very powerful tool for the characterization of different properties of graphene, such as identification of the number of layers, defects, strain, etc. [1]. Here, we report on the angular distribution of the emission of the Raman signal of graphene on glass. In general, the orientation of an emitting dipole is directly encoded in the spatial distribution of the emitted light. Fourier space imaging, also known as back focal plane imaging, allows to uniquely reconstruct this orientation from emission patterns [2]. The Raman emission from graphene can be explained by assuming point-dipolar emitters, despite graphene's extended two dimensional structure. Parameter free quantitative model calculations are presented in agreement with polarization dependent measurements. We observe that the intensity of the double-resonant 2D band of graphene is maximized when the excitation and detection polarizations are parallel and minimized when they are orthogonal, whereas that of the G band is isotropic. This is expected from polarization dependent Raman measurements [3]. Furthermore, the detection efficiency in microscopic Raman measurements is extracted from calculated patterns and is in agreement with the experimental data. In conclusion, our present work suggests back focal plane imaging as useful tool for studying the emission characteristics of nanomaterials.



Fourier space imaging of Raman scattering intensities in graphene. Comparison between experimental and theoretical back focal plane emission patterns for both G and 2D Raman signals. The arrows indicate the excitation polarization.

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Ultrastable Cellulosome-Adhesion Complex Tightens Under Load

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Conversion of biomass into fermentable sugars for biofuel production currently relies on soluble free enzymes secreted by cellulolytic microorganisms. Certain bacteria, however, produce large, modular, multi-enzyme networks called cellulosomes to more efficiently degrade lignocellulose. While network assembly is enabled by protein interactions with commonplace affinities, we show that certain cell wall associated receptor-ligand interactions prove to be extremely resilient to applied force. Through single molecule force spectroscopy and steered molecular dynamics simulations, we characterized the ligand-receptor complex which tethers the cellulosome bearing bacterium Ruminococcus flavefaciens to its substrate. The complex withstood forces of 600-750 pN, representing the strongest bimolecular interaction reported to date, equivalent to half the mechanical strength of a covalent bond. Our findings indicate inter-domain stabilization of the complex and suggest that certain network components serve as mechanical effectors for maintaining network integrity. This detailed understanding of cellulosomal network components could help in the future development of biocatalysts for production of fuels, chemicals, and pharmaceuticals from renewable plant-derivedbiomass.

Poster Presentations

Energy transfer involving core and surface states of carbon dots

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Carbon dots^(1,2) (CDs) have attracted fast growing interest due to their exceptional advantages such as high fluorescence quantum yield, chemical stability, biocompatibility, and low toxicity. Recently the fluorescent CDs have been used for bioimaging, photocatalysis, photovoltaics, as light-emitting diodes, and for lasing. However, due to the complex structure of CDs, the intrinsic mechanism and origin of the fluorescence in CDs have not yet been completely understood. Herein, by utilizing static and transient fluorescence spectroscopy, the energy transfer mechanisms in fluorescence of CDs comes from two types of states: core states, located in the carbon sp2 domains inside the CDs, and surface states, located at passivated sites mainly at the surface of the CDs. After excitation of an electron-hole pair in a core-state the energy can either decay radiatively, or can transfer to a surface state and then decay radiatively, albeit at a lower energy. Gradually shifting the excitation wavelength leads to an abrupt change in the emission wavelength, as the surface states can also be directly excited for wavelengths longer than 450 nm. These results provide insight into the internal structure of the CDs, in particular of their light emitting components, and will also help in applying these structures in wavelength-tunable nanolasers or light-emitting diode.

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Kramers Escape Problem for Self-Propelled Particles

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In the poster presentation, the dynamics of an active Brownian particle in an attractive harmonic potential is investigated. Within a 1D constant speed modeling of the particle's dynamics we consider the stationary probability distribution using both numerical and analytical approaches. By means of the results obtained hereby, subsequently the escape problem of a self-propelled particle from a metastable potential well is considered, which is discussed in detail in Ref. [1] for the case of a vanishing self-propulsion. Regarding the properties of the stationary state in a harmonic potential, two major timescales are found, governing each the translational and the rotational dynamics of the particle. Here, the particle radius is identified to be the essential quantity regulating the ratio between those timescales. For very small and very large particle radii, as well as for weak propulsion forces, approximate analytic expressions for the stationary probability distribution of the particle are derived. These analytic results compare favorably with exact numeric outcomes.Moreover, with respect to Kramers' escape problem, the analytical approximations prove to be quite beneficial, since—within their respective range of validity—the thereby obtained expressions for the escape rate coincide well with numerical findings.

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A spectral method for coupled diffusion and gene regulation

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The first step in the early development of an embryo is the establishment of spatial patterns of transcription factors which dictate the differentiation of cells in later stages of development. The concentration of transcription factors at cell nuclei during the development of a *Drosophila melanogaster* embryo is determined by coupled diffusion and gene regulation processes. Due to the small copy numbers of transcription factors and the random nature of reactions this morphogenesis is intrinsically stochastic. Conventionally, such a stochastic dynamics is studied by sampling from a large number of numerical simulations based on drawing pseudo-random numbers. However, for space dependent systems these methods are computationally very costly. Here we develop a method that allows inferring solutions of stochastic gene regulation cascades in a discretized space. Specifically, instead of only considering a single cell nucleus we are here interested in the behavior of many cells which are diffusively coupled. On the contrary to conventional methods the developed procedure follows an approach which solves for the probability distribution directly without sampling. Employing methods from quantum mechanics the master equation is expanded in its eigenbasis. A mapping scheme makes it possible to also solve for spatial degrees of freedom. For a system where the first gene regulates itself and also regulates the second gene at two cell nuclei coupled by diffusion of one species the spectral method has proven fruitful. This system could help the further in-depth investigation of the Bicoid-Hunchback interaction in the early embryo of *Drosophila*.

Nanostructuring for new perovskite solar cell absorber materials

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Alkylammonium metal trihalide perovskite absorbers have garnered a considerable amount of interest recently in the photovoltaics community. Devices fabricated from these materials achieve very high power conversion efficiencies, already superior to amorphous Si. Although very efficient devices based on planar heterojunction architecture, where the perovskite film is sandwiched between charge selective contacts, have been recently reported,^[1] the most efficient devices still require a mesoporous charge extraction layer. This is often attributed to an insufficient charge diffusion length and therefore incomplete extraction of the photogenerated charges. New perovskite materials - prepared by substituting either the halide or the metal atom - often exhibit charge collection limitations. This can be overcome by using a nano- or mesostructured titania scaffold as electron collecting anode, where the distance that needs to be travelled by the charge carriers is then reduced to the pore size of the scaffold - typically a few 10 nanometers. This could be of special importance when looking for a substitute of the toxic lead component. We propose that for new perovskite absorber material screening, it will be necessary to first incorporate them in a porous scaffold and step by step optimize the system until the charge transport properties are sufficient and there is no need for a structured electron selective layer. For this purpose we designed custom tunable titania nanostructures with pore sizes between a few 10 and hundreds of nanometers, depending on the charge carrier diffusion length in the perovskite material to be optimized.^[2]

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Figure 1: left: Titania structured with cellulose nanocrystals (pore size 8 – 15 nm); right: inverse opal structure of titania, templated with PMMA spheres (pore size: 20 – 200 nm).

Detection of the Plasmonic Circular Dichroism Signal of Chiral Molecules Self-Assembled in a Plasmonic Hotspot of Gold Nanoparticles by DNA-Origami

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We designed and implemented a DNA-based plasmonic nanostructure that can pick up the circular dichroism (CD) signal of chiral molecules located in a focus point between two plasmonic nanoparticles. The detection of a CD signal from chiral molecules in the plasmonic focus points of metal nanoparticle aggregates and on metal surfaces has been described theoretically and demonstrated experimentally^{1, 2, 3}. With the goal to observe a plasmonic CD for various chiral molecules, we designed DNAorigami-structures to bring a chiral molecule in the plasmonic hotspot of two gold nanoparticles. Plasmonic DNA origami antennas can be used for Surface Enhanced Raman Spectroscopy measurements^{4, 5}. By modifying the DNA origami structures employed in these experiments, we built a variety of addressable plasmonic hotspots. The first DNA origami structures consist of two- or three-layered square lattice DNA origami sheets with an aperture in the middle for the attachment of chiral target molecules. Single-stranded DNA extensions for the attachment of gold nanoparticles are positioned around the aperture on both sides of the origami sheet. Gold nanoparticles with diameters of 10 to 40 nanometers carrying complementary DNA sequences to the extension sequences are conjugated to the pre-defined binding sites on both sides of the aperture. In initial experiments, no aperture design was used which led to the presence of double-stranded DNA (dsDNA) in the center of the plasmonic focus point. CD signals are usually dominant around the absorption wavelength of a material – 260 nm for dsDNA. With our plasmonic particle dimer we were able to observe a CD signal in the visible region of the spectrum. In upcoming experiments, we will optimize the binding of each of the components and employ new methods to increase the concentration of the nanostructures to perform further CD measurements.

Poster Presentations

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Sequence defined oligo(ethanamino)amides for folate receptor targeted pDNA or siRNA delivery: optimization and preliminary structure-activity relationship study <u>Dongsheng He^{1,2}, Ana Krhac Levacic¹</u>, Katharina Müller¹, Petra Kos¹, Dian-Jang Lee^{1,2}, Ernst Wagner^{1,2}

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Cationic polymers are a versatile compound class with potential for the non viral delivery of different types of therapeutic nucleic acids. In order to achieve a successful and efficient delivery, just like the viral vectors, polymeric carriers also have to combine multiple functionalities. However despite their similar chemical nature, delivery of pDNA and siRNA has its own challenging obstacles which need to be addressed individually. Therefore a precise chemistry, site-specific placement of different modules and detailed structure-activity relationship studies are needed for a nucleic acid specific (e.g. pDNA or siRNA) optimization process. Herein we present sequence-defined oligomers comprising artificial polyamine domains for nucleic acid complexation, monodisperse polyethylene glycol (PEG) for surface shielding and folic acid for receptor specific cellular uptake. Via utilizing solid-phase assisted synthesis, oligomers with different topologies and additional functional domains, such as histidines for endosomal escape and tyrosines or fatty acid for the further stabilization of complexes, could be obtained. *In vitro* screening of folate receptor specific pDNA and siRNA delivery were investigated. The resulting structure activity relationships identified separate beneficial modules for pDNA or siRNA delivery. In pDNA transfections transgene expression was greatly enhanced by an increased endosomal buffer capacity, whereas siRNA delivery rather showed need for complex stabilization. In sum the work demonstrates the versatility of the presented compound platform and its potential for cargo-specific optimization.

Equilibrium Dynamics of Helical Polymers

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Biopolymers like DNA, cytoskeletal filaments or artificially designed DNA-origami fibres are famous representatives of elastic nano particles that form a helical configuration at their ground state. The handedness, radius and pitch of these helices are determined by their curvature κ and torsion τ . The statistical mechanics of such filaments is described by the helical wormlike chain model, where κ , τ are identified as the filaments' intrinsic bending and twisting rates, respectively. Here we employ Brownian dynamics simulations to investigate the thermal end-to-end distance fluctuations. We find that $<\delta R^2$ (t)> exhibits a rich scaling behavior with varying κ and τ . For $\kappa = 0$ the initial relaxation resembles the t 3/4-scaling law

as predicted by semiflexible polymer theory. In contrast, helices with a low ascending pitch angle, i. e. $\kappa > \tau$, show power law exponents exceeding 3/4 due to the additional elastic modes of the spring-like polymer conformation. The crossover region with $\kappa < \tau$ reveals a sudden intermediate relaxation regime with a scaling exponent well below 3/4. With rising τ this domain only slowly converges towards the semiflexible limiting case. Our findings demonstrate the intriguing influence of helical parameters on the dynamics of single polymer systems and can in principle help to determine structural details beyond the resolution of (static) experimental techniques.

OER catalysis in tin-doped hematite photoanodes for water splitting

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Direct utilization of sunlight for splitting water into hydrogen and oxygen is expected to be an important pillar of a future economy based largely on renewable energy. It combines harnessing the world's largest and most long-lived power source with the advantages of hydrogen as a storage medium, such as the ability to store large amounts of energy and universal usability in electricity generation as well as in chemical processes. Photoelectrochemical (PEC) water splitting at n type hematite (α -Fe₂O₃) photoanodes is appealing because of this material's abundance, chemical stability, non-toxicity and band gap suitable for absorbing a large fraction of the solar spectrum. However, its performance is severely compromised by a short hole diffusion length as well as sluggish water oxidation kinetics at the surface. Solutions to these problems are pursued by doping hematite with other metal ions, among them Sn⁴⁺, which has led to improved water oxidation performance.¹² While most previ-

ous studies attribute this to increased bulk, we show herein that enhanced catalytic activity of the water oxidation is a more plausible explanation. We found that in the mixed oxide synthesized by a sol-gel method developed in our group, tin is preferentially incorporated close to the surface in a core-shell particle structure. Intensity-modulated photocurrent spectroscopy (IMPS) and photocurrent transient analysis showed that tin doping increases the rate constant of interfacial charge transfer, and thus the water oxidation. The highest photocurrents were observed at a doping level of 20 at% tin in the synthesis solution. Furthermore, we showed that this doping also improves the electron collection efficiency for long electron pathways. This study shows that the working mechanisms of dopants in hematite can be complex and require detailed studies on well-defined model systems. For this purpose, photoelectrodes are now being prepared in our group by atomic layer deposition (ALD).

Line scan

Bulk properties

20

Surface properties

15

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30

25

20

15

10

Ó

Sn/(Sn+Fe) / a.u

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10

position / nm

Fig. 2: Radial TEM EDX line scan across Fe₂O₃ particle.



Fig. 1: J-V-Curves of Fe_2O_3 with different amounts of Sn added to the precursor solution.

Lipid-like nanoparticles for mRNA delivery

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Ethris SNIM® RNA-Technology represents a novel nucleic acid therapy platform based on stabilized non-immunogenic messenger RNA (mRNA). In contrast to commonly used mRNA, SNIM® RNA can be administered repeatedly leading to sustained expression of therapeutically active proteins even at low doses. This can be used in the therapy of hereditary diseases, as an alternative to protein therapy, and in regenerative medicine. However, the mRNA's susceptibility to enzymatic degradation in cells and its negative charge create a need of efficient delivery systems for in vivo applications. A promising approach is the complexation of the SNIM® RNA in the lipidoid formulation. Incorporation of the cationic lipid in the formulation enables stable binding of the negatively charged molecule of mRNA. The lipidoid structure is composed of an oligoamine head group modified with C12 hydrocarbon tails. The screening of different head groups in vitro indicated that the alternating structure is the determining factor in terms of transfection efficiency. In three tested cell lines: NIH-3T3 (murine fibroblasts), HepG2 (human liver carcinoma) and A549 (human lung carcinoma) the trend remained the same. We were able to proof that one reason of dramatic differences in activity between the slightly varying lipid-like structures is the buffer capacity. The acid dissociation constant in the range of pH 6.2 and 6.5 is the characteristic enabling lipid-like carriers an effective endosomal escape and, hence, efficient cell transfection. The relatively high expression of the reporter gene (Firefly luciferase) in mice after intravenous administration confirmed the utility of lipoplexes in mRNA delivery *in vivo*.

High-sensitivity measurement of DNA-Protein binding energies by automated fluorescence microscopy

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Transcription Factors that bind target DNA sequences are essential for the proper regulation of gene expression. The understanding of such transcription control networks is difficult and requires comprehensive and accurate information about the transcription factor binding affinities. Unfortunately, such data are often not available, as existing experimental methods for measuring binding preferences are low throughput or biased towards high-affinity sites. We developed a method based on Fluorescence Anisotropy (FA) to determine DNA-protein affinity landscapes in solution at large scale and with ultra-high sensitivity. We adapted a standard widefield fluorescent microscope for fully automated measurement of FA in 96- or 386-well plate format, with one affinity titration per single-well. This is achieved through a controlled delivery system, where the protein is incorporated within a porous gel matrix and the DNA-ligand diffuses through the matrix during measurements. Using a competitive binding assay, we can measure the concentration of active protein and KDs into the sub-nanomolar range with high reproducibility. The approach successfully combines automation with low sample consumption and high accuracy. We applied this technique to determine the binding specificities of 20 transcription factors participating in the well-characterized segmentation gene network of the early *Drosophila* embryo. The resulting position weight matrices show subtle but significant differences from existing ones; we evaluate their performance in a thermodynamic model¹ predicting the expression of segmentation enhancers.

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Nanoscale arrangements in soft matter films revealed by IR-SNOM

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Local ordering motives influence functional properties of soft matter systems to a large extent. In the case of proteins, folding motives such as helical are structure defining. In the case of organic thin films, molecular packing promotes the overlap of pi-orbitals and charge transfer. Since molecular vibrations are sensitive to local ordering, IR spectroscopy is a versatile technique to study local order in soft matter films; however, due to the inherently long IR wavelength, standard IR techniques lack lateral resolution in the nm range. In combination with near field techniques,¹ we are able to do nano-imaging and IR spectroscopy on soft matter films far beyond the diffraction limit, i.e. we are resolution limited only by the radius of an AFM tip (approx. 20 nm) illuminated an IR laser (s-SNOM setup by Neaspec). First, we use this technique to study polymorphism in highly ordered thin films of small aromatic molecules for organic electronics. For pentacene, nano-imaging was successfully accomplished and the coexistence of two sub-micron phases was experimentally verified.² Due to shifts of vibrational resonances, the s-SNOM is able to differentiate the distribution of bulk phase and thin film phase of pentacene with a spatial resolution of about 20 nm. Since pentacene is used for the fabrication of thin film transistors (TFTs), the identification of the local phase arrangement is of high interest since it may explain variations of charge mobility in TFTs. DNA-origami structures are another example of highly ordered molecular constructs. In collaboration with the group of T. Liedl, we want to image the structural order of DNA origami adsorbed to flat surfaces in future work.

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Radio frequency acousto-mechanical tuning of a photonic molecule

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We propose and demonstrate a dynamically tunable photonic molecule (PM) formed by two L3 defect cavities defined in a GaAs two-dimensional photonic crystal membrane. To tune the optical resonances of the two cavities of the PM we employ the acousto-mechanical deformation induced by an 800 MHz surface acoustic wave (SAW) [1]. For this frequency the cavity separation matches half the acoustic wavelength. Thus, the resulting SAW-driven sinusoidal spectral modulations of the two modes are opposite in phase giving rise to a time-dependent detuning. We monitor the mode spectrum in the time-domain by probing the time-resolved stroboscopic PL emission [2] of off-resonantly coupled quantum dots. For a PM with finite detuning, we observe for low acoustic amplitudes the established but out-of-phase spectral oscillation of two uncoupled modes. As we increase acoustic amplitudes, we resolve a characteristic anti-crossing of the two cavity modes in the time domain. From the observed splitting we are able to deduce the PM coupling strength of 180 GHz. The coupling is further confirmed in spatially and time-resolved PL maps showing delocalization of the modes over both cavities at resonance.

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Regulation of nanoparticle diffusion in the vitreous humor

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The diffusive behavior of nanoparticles in biological hydrogels is determined by the characteristics of both the gel and the particles. Hydrophobic and/or electrostatic interactions can affect the particle diffusion to a great extent. One example for such a biological hydrogel is the vitreous humor. High intravitreal mobility of pharmaceutical drugs is crucial for effective treatment of several ocular diseases. In this study we investigate the diffusive behavior of nanoparticles in ovine vitreous humor compared to porcine and bovine vitreous humor using single particle tracking. We tune the nanoparticle diffusion in the ovine vitreous by enzymatic modifications and inducing Debye screening. By investigating the particular properties of nanoparticles and hydrogel components which are involved in regulating nanoparticle diffusion, the way can be paved for the development of efficient drug delivery systems in biological hydrogels, such as the vitreous humor.

Heat driven selection of nucleic acids

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Environments shape living systems by selection. This also holds true for the first molecular replicators that had to stabilize against the equilibrium push towards dilution and decay. Especially problematic for the origin of life is the kinetic tendency of competing replicators to generate ever shorter genomes. We experimentally demonstrate that a basic thermal non-equilibrium between two geological heat reservoirs can accumulate replicating oligonucleotides and select them for increasing genome length. The interplay of molecular thermophoresis and laminar convection is able to actively filter exclusively long oligonucleotides from a steady flow through this open compartment. Trapped long strands are at the same time exposed to convective temperature cycling, which allows for exponential replication by overcoming template inhibition. Two-fold shorter strands die out in the same setting as they are simply flushed out of the system. The combination of length selection and thermal cycling renders these out-of-equilibrium compartments suitable long term habitats for the evolution of replicating oligonucleotide populations towards increasing molecular complexity. Besides the origin life, this mechanism might facilitate SELEX experiments (systematic evolution of ligands by exponential enrichment), where aptamers are selected from oligonucleotide libaries. Only the bound aptamers will remain in the trap and undergo temperature oscillations, whereas the unbound bulk will be flushed out. While standard SELEX methods consist of many time-consuming steps, here the procedure can be performed in one setting.



Heat-driven filter selecting for increased genome length. An asymmetrically heated open pore at top and bottom allows to select oligonucleotides from a steady upward flow through the pore. The interplay of gravity and a temperature gradient is sufficient to cyclically separate double-stranded DNA and drive exponential base-by-base replication reactions.

Infrared hyperspectral mapping of nano-composites

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Infrared spectroscopic analysis rests on vibrations which label any chemical compound with a characteristic "fingerprint" spectrum in the 3-30 µm wavelength region-intrinsically. Combining IR with AFM allows to perform IR chemical recognition routinely at 20 nm spatial resolution.^{1,2} Known as scattering-type scanning near-field optical microscopy (s-SNOM) this instrument has become commercially available from a CeNS spin-off (neaspec.com). Recently, we have enhanced s-SNOM by a coherent broadband mid-infrared illumination³ based on 100-fs pulses (thus even allowing nanoscale probing at ultrafast time resolution⁴); with this illumination, a complete IR spectrum is obtained at any sample position of the tip ("nano-FTIR"). By scanning the sample, an IR-hyperspectral line image or twodimensional image is recorded. The technique of nano-FTIR is being applied to thin-film organic conductors where it revealed coexisting structural phases⁵ (see also poster by D. Kalb et al. and talk by C. Westermeier). Nano-FTIR hyperspectral imaging is likely valuable in all fields of nanoscience as has already been been demonstrated in studies of protein,6 human bone,7 and interstellar-dust nano-composites.8

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Magnetic (Torque) Tweezers Experiments to Probe the Mechanics and Interactions of Nucleic Acids

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Magnetic Tweezers (MT) can probe single molecules using magnetic forces and torques. In the MT, a superparamagnetic bead is attached to the molecule of interest, for example double stranded (ds) DNA or a protein, while the other end is attached to the bottom of the flow cell. By controlling the position and rotation of magnets, above the flow cell, forces and torques can be applied to stretch and twist the molecules of interest. By varying the magnet configuration different fields can be applied for specific purposes. A "conventional" horizontal magnet configuration is suitable to analyze the forceextension behavior or to observe DNA supercoiling while rotating the molecules. Using cylindric magnets in combination with a smaller side magnet, called Magnetic Torque Tweezers (MTT), offer the opportunity to measure torque of dsDNA. As an advantage compared to other single- molecule force manipulation techniques, for a given magnet position, the applied forces in the magnetic tweezers are constant, such that no feedback is required to apply constant forces even over long

periods of time. Forces in a range of 0.1 pN up to 100 pN can be reached. In particular also smaller forces (< 1 pN), characteristic of non- covalent macromolecular interactions, can readily be applied and measured. Furthermore, our MT setup is able to track multiple beads (currently up to ~20) at the same time, enabling the collection of statistics in a single measurement run. Currently, we are investigating the precise response of dsDNA to applied forces and torques at varying salt concentrations. Preliminary analysis of force-extension and torque-rotation measurements suggest that the torsional stiffness of DNA does not depend onsalt concentration, in contrastto the bending persistence length. In addition, our MT setup allows us to address other biological challenges, for example to probe protein- nucleic acid interactions or the processes of folding and unfolding of proteins.

Microfluidic assisted self-assembly of folate-targeted Monomolecular siRNA-lipid Particles <u>R. Krzyszton^{1,2}</u>, B. Salem³, G. Schwake¹, C. Leonhardt¹, K. Müller⁴, E. Wagner⁴, J. O. Rädler¹

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The design of non-viral delivery vectors is crucial for efficient gene therapy. For optimal transport within blood and extracellular matrix used nanocarriers need to be small and stable *in vivo*. Furthermore the molecular targeting into diseased cells, facilitated by introduction of cellular receptors ligands on the surface of tailored vectors, extend their specific uptake efficiency and helps to avoid uncontrolled gene release within healthy tissues. In previous work we presented the monomolecular-siRNA lipid particles (mNALP) can be form via self-assembly during solvent exchange process. However self-assembly by this method critically depends on the mixing kinetics of the solvents isopropanol and water. Here we systematically study the self-assembly of siRNA with lipids by various mixing methods, including hydrodynamic focusing in microfluidic t-junction chip. We show that physicochemical parameters like homogeneity and siRNA encapsulation turn out to be more reproducible in microfluidic devices. We furthermore investigate the stability of mNALPs in blood serum and demonstrate that addition of small amounts of folated PEG-lipid to mNALP's results in specific binding to KB cells that overexpress the folate receptor.

Combination of a polymer-based transfection system with modified mRNA to enhance transfection *in vitro*

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Transcript therapy is based on the delivery of messenger RNA (mRNA) instead of DNA into the cytoplasm of target cells. Treating inherited genetic disorders with therapeutic mRNA is a promising alternative to DNA therapies since mRNA is translated in the cytoplasm and is only present for a period of time before being degraded. In order to create an efficient system for transfecting cells with mRNA the components of this system, vector and cargo, need to be optimized. In recent years a lot of work has been done regarding the development of non-viral polymerbased vectors, so called polyplexes, as well as the improvement of mRNA with respects to stability, expression and immunogenicity. In the present study we looked into possible synergistic effects on transfection potential and expression levels over time in NIH3T3 cells (murine fibroblasts). We combined nontoxic, poly(acrylic acid)-based vectors containing alternating pendant groups with modified mRNAs based on the SNIM® (stabilized non-immunogenic messenger RNA) technology containing a distinct combinatorial set of 5'- and/or 3'-untranslated region introducing stabilizing elements to the mRNA. Depending on the chemical structure of the pendant group cytotoxicity of the polymeric formulations was found to be lower when compared to branched poly(ethylenimine). Also all tested mRNA constructs with the differently combined UTRs showed higher expression results than the control mRNA containing no UTRs. We were not able to find a specific system comprising of a polymer and an UTR-modified mRNA that showed synergistic effects on transfection efficiency and prolongation of protein expression. Nonetheless the best performing polymer remained the most active carrier independently on the delivered mRNA construct. In addition to that each UTR-mRNA construct showed sustained expression when transfected with the polymeric formulations compared to the results obtained by transfection with DreamFect Gold™.

A Thermal, Autonomous Replicator Made from Transfer RNA

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Evolving systems rely on the storage and replication of genetic information. To this end, modern biology employs an RNA-dominated machinery to encode proteins, which in turn replicate genetic information. Before this interlinked machinery evolved, early life most probably replicated genes using a pool of short RNA sequences¹.Here, we present an autonomous, purely thermally driven replication mechanism. Instead of the chemical base-by-base replication advocated by RNA world approaches^{2–5}, the presented mechanism operates on successions of multi-base codons. It consists of a pool of RNA molecules that encodes and replicates a repetitive four-letter code. The RNAs are derived from transfer RNA (tRNA), each containing the anticodon framed by two hairpin loops. The anticodons serve as sequence-encoding toeholds while the hairpins are pairwise complementary, allowing the strands to bind sequentially. To explore the concept, we have recently shown a reduced system to work⁶. It is built of four halves of tRNA, each strand forming a single hairpin with a toehold, and replicates a two-letter code. For replication, binding energy is initially stored by thermally quenching the tRNA molecules into monom-

Poster Presentations

lecular hairpin states. Subsequent temperature oscillations connect the hairpins in a highly specific cross-catalytic reaction with a duplication time of 30 s, significantly faster than the untemplated reaction. Reaction kinetics were measured using a real time fluorescence signal and in good agreement with our kinetic model of the reaction network. In a sense, this is a physically driven version of the chemical ligation chain reaction⁷ that replicates a two-letter code cross-catalytically^{8–10}. Instead of chemical backbone ligation, matching strands are joined by physical base pairing. The approach is compatible with hydrothermal molecule traps^{11–13} and thermal microconvection¹⁴, and might have been selected for by asymmetric hydrolysis of the strand backbone¹⁵.

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Nanosized Polyplexes with Dual-Functional MTX Ligand for Enhanced Combined Cytotoxicity with Therapeutic Eg5 siRNA

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PURPOSE: The novel strategy that synthetic small interfering RNA (siRNA) can invoke RNAi responses is expected to be an excellent option for treating many incurable diseases such as cancer. However, efficient tissue-specific delivery of siRNA remains the major limitation in the development of RNAi therapy. By solid phase supported (SPS) synthesis, we have synthesized a series of sequence-defined polymers which include a cationic (oligoethanamino)amide core, cysteines (as bioreversible disulfide-forming units), and polyethylene glycol chain (for shielding surface charges) coupled to a terminal ligand. To recognize the target cells, the antifolate drug methotrexate (MTX) is employed as both targeting ligand for folate receptor-mediated uptake and as an anticancer therapeutic as it is toxic to the target cells by blocking de novo thymidylate and purine synthesis and consequently DNA and RNA synthesis. Besides, we exploit a therapeutic siRNA against eglin 5 (Eg5) that blocks mitosis to cause death of rapidly dividing cancer cells. Together we formulate Eg5 siRNA and MTX-conjugated polymer as a nanosized targeted siRNA polyplex with synergic antitumor effect.

METHODS: Size of siRNA polyplex was recorded by atomic force microscope and fluorescence correlation spectroscopy. Uptake was determined by flow cytometry and knockdown of a luciferase reporter gene was used to monitor gene silencing efficiency. To evaluate the efficacy of Eg5 siRNA, we measured toxicity by cell viability assay, mRNA expression by qRT-PCR, and aster formation of cellular DNA. RESULTS: siRNA polyplexes were uniform nanoparticles with 5.4 nm of hydrodynamic diameter. These polyplexes were taken up by KB cells in a MTX-dependent manner, and this attributed to association with folate receptor. Transfections induced significant silencing of luciferase expression in KB-eGFPLuc cells. Treatments with MTX-conjugated polyplexes containing Eg5 siRNA in KB cells triggered knockdown of Eg5, resulted in typical aster formation, and caused enhanced cytotoxicity.

CONCLUSIONS: We developed a specific and efficient siRNA carrier system with dual-functional ligand for cellular delivery and antitumor effect. siRNA polyplexes successfully delivered cargo into the target cells mainly via folate receptor. When combined with therapeutic Eg5 siRNA and MTX, the siRNA polyplexes carried out synergistic cytotoxic activity. This highly functionalized and molecule-defined carrier system for siRNA delivery could be a potential strategy for RNAi-based cancer therapeutics.

POSTER ABSTRACTS - SESSION II (LI-ZORN)

Optical printing and injection of gold nanoparticles into living cells

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The active delivery of nanoscopic objects to the surface and interior of living cells with light offers promising prospects for the development of novel molecular delivery strategies or intracellular biosensor applications. Here, we show that single gold nanoparticles from solution can be patterned on the surface of living cells with a 532 nm continuous wave laser beam by means of optical forces. Furthermore, we demonstrate that in a second step the particles can be injected into the cell by focusing the laser directly on the printed particles through a combination of plasmonic heating and the formation of nanobubbles. We find that short exposure times (within one second) are sufficient to perforate the cell membrane and inject the particles into the cell. Gold nanoparticles can be injected with a cell survival rate of > 70% depending on the laser energy which was revealed by cell viability tests.

Polarimetry of Exciton Landscapes in Monolayer MoS₂

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Monolayer transition metal dichalcogenides such as molybdenum disulfide (MoS₂) belong to a new class of truly two-dimensional materials with unique physical properties^[1]. In contrast to gapless graphene monolayer MoS₂ is a direct band gap semiconductor with optical transitions in the visible^[2] viable for potential applications in ultra-thin photosensitive electronic devices^[3]. Furthermore, coupled spin and valley degrees of freedom offer a photonic approach to valleytronic applications^[4]. We present the results of our optical studies of monolayer MoS₂ at cryogenic temperatures. In our experiments we mapped out the spatial distribution of neutral and charged excitons in large-scale monolayer MoS₂ flakes and identified with polarization resolving spectroscopy the relationship between exciton spin and valley degrees of freedom. [1] K. S. Novoselov et al., Proc. Natl. Acad. Sci. U.S.A. 102, 10451 (2005)

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High stress levels lead to transition from heterogeneous timing to synchronized cellular response of the *E.coli* Colicin E2 operon

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The production of bacteriocins in response to worsening environmental conditions is one means of bacteria to outcompete other microorganisms. Colicins, one class of bacteriocins in *Escherichia coli*, are effective against closely related Enterobacteriaceae. In a single cell study, we analyze the heterogeneous gene expression of Colicin E2, expressed from the SOS inducible *E. coli* Colicin E2 operon. We quantitatively study expression dynamics of the Colicin E2 operon in *E. coli* using fluorescence time-lapse microscopy. Different fluorescence reporter proteins allow us to observe heterogeneity in Colicin production and Colicin E2 operon and the cells response times depend thereby on the strength of the applied stress signal. While at low exogenous stress levels all cells eventually respond after a given time (heterogeneous timing), high stress levels lead to a synchronized stress response of all cells about 75 min after induction via stress. A heterogeneous response in combination with heterogeneous timing can be biologically significant. It might enable a bacterial population to endure low stress levels, while at high stress levels an immediate and synchronized response may allow elimination of closely related bacteria competing for resources. Furthermore we could demonstrate, that the amount of Colicin released is dependent on *cel* (lysis) gene expression. Future investigations will focus on transcriptional as well as post-transcriptional regulation affecting the dynamics of Colicin expression and release. A separation of powers - force and function of the versatile cohesin-dockerin interaction <u>Lukas F. Milles</u>, Wolfgang Ott, Ellis Durner, Markus A. Jobst, Klara H. Malinowska, Constantin Schöler, Tobias Verdorfer, Michael A. Nash, Hermann E. Gaub

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How can we assemble protein networks? The answer to that question may have have been discovered with the cellulosome, a modular system of numerous enzymes expressed by a multitude of thermo- and mesophilic bacteria, most prominently *C. thermocellum*. It excels at breaking down lignocellulosic biomass through hierarchical assembly. The modular basis of cellulosome network formation is the high affinity cohesin-dockerin receptor-ligand interaction. Different bacteria produce structurally different cellulosomes but exhibit homologous cohesin-dockerin pairs. We present a sampling of representative interactions, characterize their affinities and find them to be distinct to their position in the cellulosome network. Single-molecule force spectroscopy is used to measure the force necessary to separate a cohesin-dockerin complex. The resulting force-extension traces show various unfolding pathways with a recurring fingerprint. We developed a classification algorithm to determine the most probable unfolding pathway clusters through dimensionality reduction techniques. Furthermore we demonstrate the suitability of a cohesin-dockerin system as a tag to study enzyme unfolding pathways in single molecule pulling experiments. The cellulosome's constituents hold the potential to become versatile connectors for nanoscopic protein assembly and have already proven their aptitude as reliable tags for single molecule force spectroscopy experiments.

Polymerization and protection of nucleic acids in a prebiotic environment

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The formation of the first replicating systems capable of Darwinian evolution in an aqueous environment was likely a crucial step towards the evolution of life. Current opinions on the prebiotically plausible synthesis of single nucleotides however require multiple steps of drying and UV irradiation^[1]. We therefore propose porous rock at hot springs near sea level as sites where both dry and wet conditions can be combined. A temporarily dry surface of the rock would allow reactions where the synthesis and polymerization of nucleotides can take place^[1,2]. Washing these products into the pores would then enable their accumulation and further polymerization by harnessing a temperature gradient across the pore^[3]. This can lead up to a point where the polymers form macroscopic structures that are even better protected against dilution. In our experiments, we show a dry polymerization of 3',5'-cyclic GMP to long nucleic acid strands without the help of catalysts or enzymes at moderate temperatures^[2]. In addition, we

demonstrate the enzyme-free compaction of two-base (G & C) nucleic acid strands in a simple temperature gradient across a water-filled pore. The strands are capable of polymerization and branching via hybridization of complementary parts and form millimeter-sized structures that are stable against dilution and small water flows. This also paves the way for a selection mechanism in which only polymerizing strands are retained in the system, while others are flushed out over time. Our goal is to eventually combine dry and liquid states in one pore in order to increase the possibilities of reaction pathways in the same setting, while maintaining the advantages of a small and separated compartment.

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Gelation of nucleic acids. Two-base 24mers were accumulated for ~100 min. The strands consist of three self-complementary parts that allow them to form a network. This results in a macroscopic gel that is stable against water flows and can survive for days without active accumulation. Strands were labeled with SYBR Green I in 1x PBS buffer. Scale bar: 100µm.

Chemiluminescence of an anthracene-based UiO-68

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Numerous possible combinations of metals and organic linkers that comprise metal-organic frameworks (MOFs) offer great topological and chemical diversity within those strucutres. These features make them ideal candidates for applications in gas storage, catalysis, and chemical sensing.^[1] For chemical sensing applications, luminescence represents a powerful read-out concept since the measurements are usually very sensitive and offer high resolution, with even single molecule visualization. The distinct advantage of chemiluminescence



in comparison to fluorescence is the absence of external light sources. Thus implies that the analytically relevant emission can be measured against a completely dark background, resulting in a high sensitivity of this transducer concept. In the present work, we developed a derivative of 9,10-diphenylanthracene, which is known to have chemiluminescence properties. We synthesized an UiO 68-related structure with this building block, resulting in a MOF with large pores and high surface area.^[2] Strikingly, the UiO-68(anthracene) MOF shows strong chemiluminescence once exposed to a solution of bis(2,4,6-trichlorophenyl)oxalate and hydrogen peroxide. We anticipate that the implementation of a chemiluminescent transducer concept within MOFs will greatly extend the scope of MOF-based sensors.

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Mechanics of dimeric von Willebrand Factor under varied pH conditions probed by Single Molecule Force Spectroscopy (AFM)

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Von Willebrand Factor (VWF) is a giant multimeric blood glycoprotein that senses shear flow irregularities in the blood stream. At sites of vascular injury, where shear forces are increased and of elongational character, VWF extends and subsequently promotes platelet adhesion^[1, 2]. Since formation of a platelet plug is essential for hemostasis, defects in or deficiency of VWF can lead to severe bleeding disorders, such as found in von Willebrand disease. Understanding the structural rearrangements within VWF when stretched in the blood stream is of fundamental interest from both a biophysical and medical perspective. Using AFM-based Single Molecule Force Spectroscopy, we probe the mechanical signature of VWF dimers, the smallest repeating subunits of VWF. Small peptide tags at opposite termini of the dimers allow pulling specifically and in a well-defined geometry. Varied pH conditions help to interpret the obtained force-extension profiles, as pH was shown to influence the static structure of VWF

dimers and thus facilitate their correct multimerization in the endothelial Golgi apparatus ^[3]. From a more medical perspective, acidic conditions are found at sites of inflammation in the blood. Overall, we report on the mechanics of dimeric VWF under physiologically relevant pH conditions.

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Conductivity in Polymeric Electrode for Roll-to-Roll Printed Organic Electronics: Evolution of Nanostructure and Molecular Orientation in PEDOT:PSS

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Since the invention of organic electronics (OE), organic thin films have attracted immense interest among researchers from all over the world. The transparency and the potential of manufacturing of OE devices on flexible substrates enables numerous smart design opportunities as well as roll-to-roll (R2R) production. The performance of optimized OE devices strongly depends on the optimum length scales of the nanomorphology. In this work, the underlying morphological changes in the film volume are probed with ex-situ and insitu X-ray scattering techniques: grazing incidence small and wide angle X-ray scattering (GISAXS and GIWAXS).[1] The brittleness of the commonly used indium tin oxide (ITO) is one of the main limiting factors for the R2R production of OEs. Flexibility however, is one of the main advantages of OE devices over conventional electronics. In this context, we show that the addition of co-solvents increases the conductivity of PEDOT:PSS polymer blend films by more than three orders of magnitude, making it suitable as an ITO replacement. The underlying structural evolution is probed with GISAXS and GIWAXS (DESY, Hamburg). As a result, the enhancement in conductivity is ascribed to fundamental morphological changes and for the first time to molecular reorientations within crystalline domains. [2]

In order to realize the goal of up-scaling, techniques such as printing are needed. Film formation typically takes place within a couple of seconds and involves complicated structure and morphology evolutions. Hence, to understand the formation of printed films, in-situ investigations are necessary. We realize the printing of PEDOT:PSS films with a self-made slot die coater, specially designed for the implementation at a synchrotron facility (ALS/LBNL Berkeley & UMass Amherst) for in-situ GIWAXS measurements. This experiment provides a comprehensive understanding of the nanomorphology evolution for directed device optimization.^[3]

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Figure 1: Ex-situ and in-situ GISAXS and GIWAXS investigations of PEDOT:PSS polymeric electrodes. The enhanced conductivity of PEDOT:PSS by co-solvents is ascribed to improved morphology and molecular orientation in crystalline domains.

Identification of siloxane mechanochemistry with single molecule force spectroscopy and ab initio simulations

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The mechanical stability and the degradation of individual chemical bonds can be investigated by single molecule force spectroscopy. Quantum chemical calculations of possible reactions including effects of external forces aid the interpretation of these experiments. We have investigated the force-dependent lifetime of several chemical reactions using the atomic force microscope (AFM) in force-clamp mode. Single molecules of carboxymethylated amylose (CMA) were coupled with different silanes between a glass substrate and the AFM tip.1 Individual CMA molecules were stretched with different clamp forces and temperatures until a bond scission was observed. Reaction kinetics of the experiments exhibited single or bi-exponential behaviour. Bi-exponential kinetics indicates at least two separate components in the ensemble of molecules investigated: for N1-[3-(trimethoxysilyl)-propyl]diethylenetriamine (DETA), with three ethoxy groups, and 3-aminopropyl(diethoxy)methylsilane (APDEMS), with two ethoxy groups a bi-exponential behaviour was observed, while for (3-aminopropyl)-dimethyl-ethoxysilane (APDMES), with one ethoxy group, a single exponential behaviour was observed. One ethoxy group is always bond to the surface, and via the additional ethoxy groups in DETA and APDMES, the silane can form cross-links to neighbouring silane anchors, while for APDMES cross-linking to neighbouring slianes is not possible. The bi-exponential behaviour, of DETA and APDEMS, can be explained by the fact that not at all silanes are always cross-linked. Data analysis with Arrhenius kinetic models based on a Morse potential was used to identify force independent kinetic parameters. For all three silanes one decay channel showed the same reaction parameters, which indicates a reactions at the siloxane bond, when the silane is not cross-linked. The activation energy was approximately 50 kJmol⁻¹ and the Arrhenius prefactor approximately 10⁸ s⁻¹. To explain the detail of the reaction pathway several possible chemical reactions were calculated using quantum chemical methods². To obtain force dependent kinetic parameters the External Force is Explicitly Included³ method was used. The base catalysed hydrolysis of the siloxane anchor exhibited nearly the same activation energy and force dependent kinetic parameters as in the experiment.

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Lissajous rocking ratchet in quantum dots

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Symmetries are a very important concept of physics - the most famous one being the CPT symmetry. Breaking symmetries often gives rise to interesting effects and, in particular, breaking the time-reversal symmetry is a requirement for many applications such as information processing. Here we present such a concept based on a quantum dot (QD) electrostatically defined in a AlGaAs/GaAs heterostructure. We break time-reversal symmetry by periodically modulating its barriers such that a single electron tunneling current occurs. The current direction can be controlled by introducing a phase difference between the two periodic signals. We show that our QD resembles a Lissajous ratchet. A consistent theoretical model based on scattering matrix formalism describes our experimental findings. Similar devices could be realized in a large variety of systems, for instance in nanomechanical or superconducting circuits. Possible applications include noise management, filtering and signal routing.

Facilitated diffusion mechanism of microtubule polymerases

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The diffusive motion of proteins on filamentous structures in the cell is vital for several cellular functions like gene regulation and cytoskeletal dynamics. A targeting mechanism of the respective reaction sites that is faster than the limit imposed by three dimensional diffusion is crucial in both cases. In this respect, recent experimental studies suggested that a facilitated diffusion mechanism due to reduction of dimensionality in the diffusive motion is utilized by two key players involved in regulation of microtubule dynamics, MCAK and XMAP. To quantify this facilitated diffusion mechanism on a theoretical level we investigate a stochastic lattice gas model for enzymatic activities at microtubule ends such as polymerization and depolymerization. We quantify our models using data accessible from experimental studies. This allows for a direct comparison between theory and experiment. For both proteins, we propose a model that is in excellent agreement with experimental findings. Our results show that one dimensional diffusion serves as a crucial mechanism to enhance the enzymatic activity of these proteins. We hypothesize that this mechanism ensures that regulation of microtubule dynamics by MCAK and XMAP becomes feasible under physiological conditions. Furthermore, we observe that the facilitated diffusion mechanism operates most efficiently at cellular enzyme concentrations.

Nanoscale DNA origami based plasmonic ring structures

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Nanoscopic materials for light manipulation are of great interest due to their possible applications in signal modulation, metamaterial research and optical sensing. With the aim to control electromagnetic waves at visible frequencies, we created selfassembled nanostructures with subwavelength features that exhibit collective plasmon modes in the visible region. One concept for the experimental realization of artificial optical magnetic modes relies on the arrangement of nanoparticles in a ring structure^{1,2}. The bottom-up strategy of DNA-self-assembly has been demonstrated to be a viable method to construct building blocks for plasmonic metamaterials³. Herein we applied this method for the fabrication of ring structures composed of four to eight precisely organized single metal nanoparticles by using a ring-shaped DNA origami structure of ~ 60 nm diameter. The optical response of these metamolecules was measured by bulk absorption spectroscopy as well as by single structure scattering spectroscopy. The unique plasmonic features were then compared to computational simulations. In conclusion we demonstrated the possibility to create fluids containing optically active metamolecules.

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Fig. 1: DNA-origami – AuNP ring structures. a, Plasmonic ring structures were constructed by arranging gold nanoparticles on ring-shaped DNA origami structures with specific predefined attachment points. Scale bar 40 nm. b, Wide field TEM micrograph of the DNA-origami – AuNP hybrids. Scale bar 400 nm. Inset: Zoom in on one structure. Scale bar 40 nm. c, dark-field scattering spectra of a single structure surrounded by water (black) and air (red) and corresponding numerical simulation.

Novel thermoelectric films based on polymer-nanoparticle composite

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Thermoelectric materials gained interest over the past years, due to a high interest in renewable, sustainable and environment-friendly technologies for energy conversion. In thermoelectrics, a temperature gradient is applied to both ends, which leads to a change in the charge carrier distribution due to their flow from the hot to the cold side, depending on the electrical conductivity of the material. It is feasible to have a low thermal conductivity, which prevents the propagation of phonons and heat through the material. A measure for the efficiency of a thermoelectric material is the so-called figure of merit ZT:

$$ZT = \frac{\sigma S^2}{\kappa}$$

 σ describes the electrical conductivity, S the Seebeck coefficient or thermopower and S the thermal conductivity. σS^2 is also often referred to as the power factor. So far, thermoelectric materials mostly comprised inorganic components, such as bismuth and tellurium. Although these materials show high ZT, they suffer from problems such as high energy consumption for production, or environmental concerns because of potential toxicity of the components and/or low abundance.In order to overcome these restraints, we present a hybrid approach for making thermoelectric films by combining the co-solvent-treatedpolymer blend PEDOT:PSS with inorganic nanoparticles. This strategy is especially appealing because of the high tailorability of polymer morphologies, the high availability of polymer materials and the possibility for solution-based processing at ambient conditions. In this context, we realized a functioning thermoelectric system and now aim for the increase of the ZT-values. Measurements were carried out regarding electrical conductivity, Seebeck coefficient and film thickness. In addition, first experiments were used to investigate the correlation of the distribution of nanoparticles within the film with the measured thermoelectric properties using resonant soft x-ray scattering.

Non-equilibrium transport through a QD using Keldysh-fRG 1

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The functional renormalization group (fRG) is a powerful resummation technique which recasts the computation of diagrams into a differential equation. We apply the non-equilibrium (Keldysh) version of it to a single impurity Anderson model. From previous studies it is known that the truncated fRG-scheme employed violates the particle-conservation law. Thus, the resulting value of a given observable, e.g. the conductance, depends on the precise representation used. We show the dependence of the differential conductance out of equilibrium on the representation for finite bias voltage at different temperatures and magnetic fields. In future, we hope to apply this technique to a quantum point contact (QPC). The QPC is assumed to be described by a onedimensional, inhomogenous, interacting region coupled to noninteracting leads. So far, a different implementation of fRG has been used to explore the equilibrium properties of this model. It shows striking qualitative agreement with the measured 0.7 anomaly.

Click chemistry – a versatile method for enzyme immobilization in large pore colloidal mesoporous silica nanoparticles

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Recently, we have developed several bare and modified colloidal mesoporous silica systems, for example uniform MSNs functionalized with numerous organic moieties.^{[2][4-5]} However, the synthesis of MSNs with large pores (LP_MSN) and well-defined small particle sizes is an important challenge, as the pore size of conventional silica nanoparticles (around 4 nm) is unsuitable for the adsorption of larger biomolecules such as enzymes and small interfering RNA (siRNA). The immobilization of enzymes for catalytic processes is a key application of mesoporous silica materials as the pores protect the fragile cargo from external influences such as degradation. In this contribution, we demonstrate the optimization of the pore diameter and the molecular functionalization of the mesopores in order to allow the facile diffusion and attachment of guest molecules into the host. An increase in pore diameter to values larger than 5 nm was achieved with micellar expanders, large molecular weight surfactants and novel fluorinated co surfactants.⁽⁶⁾ The employed synthesis procedure yields homogeneous azide-functionalized LP_MSN-N₃ of about 120 nm in size with ultra-large pores of 10 nm, using a direct co condensation approach to introduce the organo-functionality. The azide-moieties in the porous framework allow for a mild and biocompatible click chemistry reaction with two acetylene-functionalized enzymes of different sizes (sp cytochrome C and sp-trypsin). The catalytic activity of the immobilized enzymes was investigated with fluorogenic reactions and it was demonstrated that both enzymes are highly

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active in their immobilized state compared to previous studies and free enzymes; a 2.9 times higher activity was observed for LP_MSN-CytC compared to the free enzyme.

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Characterizing Cell Motility and Transmigration in Ring Shaped Micro Patterns

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Cell migration is important in many biological processes such as embryogenesis, wound healing, or cancer metastasis. To understand the formation of tumors and the effect of drugs, a detailed characterization of the migration behavior is important. Furthermore the ability to overcome barriers like the basement membrane is a key indicator for the aggressiveness of different cancer cells. Therefore a systematic approach for studying transmigration behavior is necessary to characterize the invasiveness. Here we study single cell migration constrained to a micropatterned ring-shaped lane. On such tracks cells perform a 1D

persistent random walk. Analyzing large arrays of these patterns in parallel, we are able to evaluate characteristic velocities and persistence times of a cell line with high accuracy. By introducing a gap of defined size and chemical composition in the ring shaped lane we study cell behavior at defined chemical interfaces. At the chemical border cells either turn around or transmigrate over the barrier. Studying the transmigration probability systematically, we find a steady decrease of transition probability with increasing barrier width. Thus, this system will allow the detailed comparison of cell lines with varying invasiveness.

Models for Angiogenesis on microstructured surfaces

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Angiogenesis, the growth and formation of novel blood vessels from preexisting vessels, is an important physiological and pathophysiological process involved in wound healing but also in cancer progression. However, dynamics of angiogenesis in general and the impact of physical factors are barely understood. We try to model different cellular processes of primary endothelial cells using micro-structured surfaces. With the help of micro contact printing we bring HUVECs (= Human Umbilical Vein Endothelial Cells) into defined cellular shape and stress conditions in order to model 3D migration, tip cell formation and fibronectin-fibrillogenesis. All these processes have been shown to play an important role in angiogenesis. We use primary endothelial cells, which are not adapted to 2D cell culture, to establish an easy accessible model system for imitating 3D migration on a flat surface. This model has been proposed before for 3T3 fibroblast, but HUVECs show some striking differences. Our results reveal that 1D migrating endothelial cells share a lot of properties compared to 3D migrating cells, regarding their overall morphology as well as their cellular response to selected small molecule inhibitors.

DNA Nanotubes as Intracellular Delivery Vehicles in vivo

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We investigated the use of DNA-based nanotubes as carrier systems for CpG delivery and their effect on immune cells *in vivo* and in real time. DNA nanotubes were produced using the single-stranded tile assembly method by mixing 48 unique DNA strands to form 8-helix nanotubes with a length of ~40 nm and a diameter of ~8 nm. Unmethylated CpG sequences are recognized by Toll-like receptor 9 (TLR9), present on lymphocytes

and antigen-presenting cells, incl. macrophages, and thus initiate an innate immune response. To test the immune stimulation *in vivo*, CpG-decorated DNA nanotubes were microinjected into the cremaster muscle of anesthetized mice. Using *in vivo* microscopy, we observed that CpG-decorated DNA nanotubes were rapidly internalized by tissue-resident macrophages and colocalized with late endosomes/lysosomes in these cells. Only microinjection of CpG-decorated DNA nanotubes but not of plain DNA nanotubes or CpG oligonucleotides induced a significant increase in leukocyte adhesion and transmigration in postcapillary venules of the cremaster muscle. Interestingly, CpGdecorated DNA nanotube-induced leukocyte recruitment was almost completely abolished in animals treated with an inhibitor of mast cell degranulation. Confocal microscopy of immunostained muscle tissue revealed that only after treatment with

CpG-decorated DNA nanotubes, nuclei of cells surrounding the microinjection site were positive for phosphorylated p65, indicating TLR-9-mediated activation of the NF- κ B pathway. Taken together, these *in vivo* findings suggest that DNA nanotubes are promising delivery vehicles to target tissue macrophages. The immunogenic potential apparently depends on the decoration of DNA tubes with CpG oligonucleotides.

Tip-enhanced Raman Spectroscopy (TERS) – Excitation Power Dependence?

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Tip-enhanced Raman spectroscopy (TERS) utilizes the enhanced electric field in the proximity of an optical antenna to achieve nanometer spatial resolution and high detection sensitivity⁽¹⁾. However, the Raman enhancement mechanism under strong local field still remains unclear. A recent report indicates a contribution from non-linear Raman scattering in the case of very strong enhancement⁽²⁾. To study this, we conduct power-dependent confocal and tip-enhanced measurements by simultaneously recording the excitation laser intensity and the corresponding Raman G-band signal of single carbon nanotubes

(CNTs) dispersed on a gold film. Here we discuss a possible contribution of stimulated Raman scattering using a single excitation laser frequency. Our results will contribute to our understanding of tip-enhanced mechanism and shed light on experimental findings.

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Cell manipulation using Surface Acoustic Waves

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Studying cell proliferation and migration is crucial not only for scientific interests, but could also be used for a medical purpose. Therefor adhesion phenomena of different cell-material combinations can be examined to determine specific properties. For an optimal implementation of materials in living environments, a thorough characterization of cell adhesion properties, both kinetics and strength, is required. Here we present a miniaturized (~ 100 μ l) lab-on-a-chip implant hybrid system which allows to quantify cell (de-) adhesion under dynamic conditions mimicking those of physiological relevance. Surface acoustic

waves on optical transparent chips are used to create a microfluidic shear spectrum ranging from 0 - 400 s⁻¹ which the cells are exposed to. We demonstrate its applicability with a model of an osseointegration study using SAOS-2 cells on medical implant material samples. The great advantage of DANI compared to present-state cell adhesion probing systems is that it requires only very few lab consumables, live observation of the cells and arbitrary material-cell combinations. Further on, the measurement chamber allows temperature and pH-value control (e.g. to generate physiological conditions).

Game Theory on the Nanoscale

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In evolutionary game theory, cyclical dominance is a simple set of rules, that can lead to spatio-temporal pattern-formation^[1]. The ability of such systems to form these patterns depends on the diffusion coefficient and the system size. The DNA-Toolbox is an well-established set of enzymes (polymerase, exonuclease, nickase) and DNA-strands, where the amplification of one DNA-species can activate or inhibit the production of another species^[2]. With a similar system, predator-prey-dynamics have already been observed^[3,4]. Linking three DNA-species together, such that A inhibits B, B inhibits C and C inhibits A, allows cyclical dominance on the nanoscale. Using a simple model^[2] and a finite element simulation software package (COMSOL), the behavior can be simulated in 0D, 1D and 2D. In the latter case, moving patterns like spiral waves build up. The three species replace each other dynamically according to their cyclical dominance.

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Microparticle formulation of mRNA for sustained protein production in cells

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Stabilized non-immunogenic messenger RNA (SNIM[®] RNA) is being established as a favourable alternative for enzyme replacement or gene therapies. SNIM[®] RNA can be administered repeatedly leading to expression of therapeutically active proteins even at low doses. In some cases, a prolonged expression is desired to achieve this goal only with a few applications. Hence, SNIM[®] RNA has been compacted and concentrated into PLGA (poly(lactic-co-glycolic)acid) microparticles of adjustable sizes to increase the local sustained release of the messenger RNA. In this work of research it is shown that a prolonged expression can be achieved by using microparticles as the delivery agent in vitro on mesenchymal stem cells compared to a traditional lipidic nanoparticle formulation of SNIM® RNA. The goal of this project is to develop a drug delivery system which can be used for local administration in different body tissues like bone, muscles or skin. The microparticles have been investigated by mRNA expression tests *in vitro*, and additionally by light – and electron microscopy for size analysis.

From Genes to Protein Mechanics on a Chip

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Mechanical forces acting on proteins play a pivotal role in biological systems. By applying forces to single molconformational changes and energetic barriecules. ers along unfolding pathways can be probed by SMFS. Since low experimental throughput has significantly limited the capacity to screen libraries of proteins, we developed a versatile microfluidic system to address these issues. Our platform enables parallelized force spectroscopy utilizing cell-free in vitro gene expression, covalent protein immobilization and subsequent measurements of mechanical properties at the single molecule level in a streamlined format with one single cantilever. A PDMS microfluidic chip on a glass slide seals engineered DNA spots to provide micro reactors for protein synthesis. Expressed fusion proteins covalently attach to the glass surface at their N-termini and display free dockerin domains at their C-termini. With a single cohesin-functionalized cantilever, unfolding pathways and unbinding characteristics of multiple different proteins can be probed. Our example library contained structural proteins, cytoskeletal constituents, enzymes, and fluorescent proteins, which we were able to detect by their specific unfolding fingerprints. Analysis of contour-length increments and rupture force - loading-rate data characterizes the constructs and are compared with computational methods. As an application of this novel system, mutant variants of individual receptor-ligand proteins can be constructed, immobilized and measured on a single molecule basis to screen for candidates in protein design. For example, saturation mutagenesis of single residues responsible for binding their counterpart can be performed and characterized on a single device. This method provides a unique means of comparing forced dissociation and unfolding pathway characteristics of engineered proteins.



Method workflow. (a) A gene array is spotted onto a glass slide and a multilayer microfluidic chip featuring 640 unit cells is aligned to the DNA microarray and bonded to the glass slide. Each unit cell comprised a DNA chamber, a protein chamber, and superseding elastomeric control valves actuated by pneumatic pressure. (b) Control valves are utilized for spatially selective surface modification of each protein chamber with PEG-CoA, for fluidic isolation of each chamber prior to in vitro expression of the microspotted DNA and for fluorescent labeling with TagRFP-Cohesin. (c) After removal of the microfluidic device, the resulting well-defined, covalently attached protein microarray are accessed from above with a functionalized AFM cantilever. Singlemolecule unfolding traces of each of the protein constructs are thus acquired sequentially at each corresponding array address with a single cantilever in a single experiment.

CeNS Workshop Venice 2014

Mesoporous silica nanoparticles as drug delivery platforms^[1]

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Nano-sized mesoporous silica particles (MSN) with high colloidal stability attract growing attention as drug delivery systems for targeted cancer treatment. These MSN offer a high pore volume, a defined and tunable pore size, and various functionalization possibilities.^(1,2) Recently, we have developed a strategy to incorporate molecular functionality at different locations of colloidal mesoporous silica nanoparticles. For targeted drug delivery, the triggered release of bioactive compounds at specific locations, times and conditions is highly desirable. Here, we present our recent studies on different capping and release mechanisms for

controlled and targeted delivery of model drugs into cancer cells. With live-cell spinning-disc fluorescence microscopy we were able to monitor the uptake of MSN by single cells, trigger endosomal escape by photoactivation and detect the release behavior from the particles (cf. Figure). The functionality of the targeting ligands which have been attached to our nanoparticle system was evaluated by performing competition experiments with free ligands. We investigated several delivery platforms based on MSN including a supported lipid bilayer as capping system, or pH-responsive gatekeepers such as polymer, enzyme or den-



dron structures. Controlled release could be demonstrated either with light activation, reduction of the pH or reductive milieu inside living cells. For demonstration of successful intracellular release of the encapsulated guest molecules we either monitored spreading of fluorescent model drugs or the destruction of fluorescently labeled cell compartments. The obtained results demonstrate that multifunctional MSN are a promising and flexible platform for drug delivery applications, such as cancer therapy.

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Schematic representation of different stages of a targeted cellular uptake of a multifunctional MSN and controlled release of the cargo into the cytoplasm of cancer cells. 1) Active docking to cell surface receptor of a nanocarrier via targeting ligands; 2) process of ligand/receptor-mediated endocytosis; 3) MSN entrapped in endosome; 4) intracellular transport and acidification of endosome; 5) triggered endosomal escape of nanocarrier, thus obtaining access to the cytoplasm; 6) controlled delivery of the cargo inside the cell. ¹¹

Poster Presentations

Conformational changes of Sti1 when Hsp70 or/and Hsp90 bind

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Heat shock proteins like Hsp70 and Hsp90 work together as chaperones to help nascent proteins reach their final functional conformation. Understanding the underlying mechanisms of chaperone assisted protein folding may thus help to prevent diseases like Alzheimer or Parkinson. The cochaperone Sti1 helps to bring the chaperones Hsp70 and Hsp90 in contact. Sti1 has five domains, two aspartate and proline rich (DP) domains and three tetratricopeptide repeat (TPR) domains and a flexible linker. The different domains form three functional elements, a flexible N-terminal module (TPR1 and DP1), a long linker region and a rigid C-terminal module (TPR2A, TPR2B

and DP2). Sti1 has one binding site for Hsp90, in the TPR2A domain, and two binding sites for Hsp70, in the TPR1 and the TPR2B domain (Figure 1a). On a solution based setup with MFD (Multi-parameter Fluorescence Detection) in combination with PIE (Pulsed Interleaved Excitation)1 we use spFRET (single pair Förster Resonance Energy Transfer) measurements to show that the two TPR2 domains are rigidly connected (Figure 1b) and the distance of the ends of the linker does not change in the presence of Hsps (Figure 1c). Furthermore, we encapsulated the protein in 200 nm large vesicles, which we then immobilized to the surface2. We could demonstrate, by using surface based



spFRET with TIR (Total Internal Reflection) excitation, that Hsp90 influences where Hsp70 binds. In the absence of Hsp90, the TPR2B domain is the preferred Hsp70 binding domain. The Sti1 molecule is in a more compact conformation (Figure 1d) and Hsp70 prefers to bind to the TPR1 domain, when Hsp90 binds. Taken together, our results suggest that the presence of Hsp90 regulates the binding of Hsp70 and assures the transfer of Hsp70 between the two modules, which is maybe responsible for the efficient handover of the client protein between Hsp70 and Hsp90.

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Double Gate Organic Thin Film Transistors on Thin Foils for Biosensing

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There is a need for cheap, flexible and lightweight biosensors. Biosensors based on organic electronics match these requirements. We have demonstrated a transducer based on an Organic Thin Film Transistor (OTFT). The organic semiconductor was encapsulated by a thin layer of tetratetracontane (TTC) to enable stable operation in electrolyte [1]. However, the bottom gate of this device has still been a silicon chip. We replaced the Si gate with a thin metal electrode and the silicon oxide dielectric with a biocompatible dielectric, parylene to fully realize the advantages of organic electronics. Thin foils of parylene with a thickness of ca. 2 μm can also serve as the substrate for the OTFT, resulting in flexible devices. The electrode structures of the OTFTs are micro patterned by photolithography. Such OTFTs can be operated in electrolytes [2]. Here we show operation of OTFTs operated as double-gate thin film transistors based on parylene as substrate and bottom gate dielectric. The encapsulation has been changed from TTC to parylene-C, since these films yield a higher capacitance and are crosslinked. Measurements are done in 10 mM Dulbecco's PBS buffer solution. The threshold voltage of the top gate can be adjusted by the applied bottom gate voltage. This allows maximising the transconductance of the OTFT. The OTFTs are mounted in a commercial plastic micro fluidic flow chamber. Six individual transistors are placed in the flow channel and can be read out in parallel via hot-switching through the individual transistors. Detection of urea is demonstrated with OTFTs functionalized with the enzyme urease. Urease catalyzes the hydrolysis of urea, which results in a pH shift. This pH shift is measured by the transistor.

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Manipulation of cell proliferation and migration using surface acoustic waves <u>Christoph Westerhausen^{1,2}</u>, Manuel Brugger^{1,2}, Melanie Stamp^{1,2}, Achim Wixforth^{1,2}

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During tissue injury large numbers of endothelial cells are injured, destroyed or removed. After hemostasis the wound needs to be sealed. Thus, the remaining cells have to rebuild the tissue by proliferation and migration to restore its function. Here, obviously, time is a critical factor. Thus, an exciting question in biological physics is: "Can you enhance cell growth?" and, considering clinical surgery: "Can you manipulate velocity and direction of cell growth? ". Beneath radiation, pressure or temperature changes to manipulate cell growth, mechanical effects are of special interest. In the past, static mechanical treatment has been shown to stimulate cell growth under specific conditions. However, highly controllable dynamic manipulation of microscopic systems respectively living cells is enabled by the technology of surface acoustic waves. Here we demonstrate the bio compatibility and bio functionality of such a bio reactor for dynamic cell manipulation. Saos2 cells on a SiO_2 covered LiNbO₃ substrate are continuously treated with acoustic waves for 72 hours. By the combination with established wound healing assays we show significantly increased proliferation respectively migration of Saos2 osteoblasts compared to reference samples. Moreover we give an outlook on continuative experiments to explain the physical reasons of the observed effects.

Synthesis of nanostructured metal oxide electrodes for electrochemical lithium insertion

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The development of electrode materials offering both high energy density and high power is urgently required in the field of electrochemical energy storage. Materials with very short charging times are highly desirable for the use in electric vehicles and mobile electronics and will enable many envisioned applications. Nanostructuring is known to be one way to drastically alter the properties of materials and therefore is considered as one of the key routes towards the improvement of already commercially applied electrode materials. Using a novel solvothermal synthesis approach in tert¬ butanol we develop new pathways for the fabrication of different ternary metal oxide nanoparticles for the application in lithium-ion batteries. Furthermore, our research focusses on their controlled assembly into continuous networks. With this method we were able to obtain fully crystalline interconnected porous frameworks composed of ultrasmall lithium titanate $Li_4Ti_5O_{12}$ (LTO) spinel nanocrystals.⁽¹⁾ These were shown to be the fastest everreported titanate morphologies as anode material for lithium insertion. We have found that the reduction in crystal size also leads to a change in bulk ion transport properties of the nanocrystals, explaining their extremely fast electrochemical lithium insertion rates. Here we show the extension of our successful synthesis strategy to the development of nanostructured thin films of lithium cobalt oxide $LiCoO_2$ (LCO), which was the first commercially used cathode material in lithium ion batteries. The reduction in size led to faster charge and discharge processes in nanostructured LCO compared to bulk material.

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Flow and diffusion in channel-guided cell migration

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Collective migration of mechanically coupled cell layers plays a notable role during embryonic development, wound healing and cancer progression. Confluent epithelial sheets are well-studied and spontaneous formations of swirls as well as glass-like dynamic arrest as a function of cell density have been uncovered. In contrast, the flow-like properties of one-sided cell-sheet expansion in confining geometries are not well understood. We studied the short- and long-term flow of Madin-Darby canine kidney (MDCK) cells as they move through microchannels, using single cell tracking and particle image velocimetry (PIV). We found that a defined stationary cell current emerges when data is averaged over characteristic spatial and temporal scales. The averaged flow-field exhibits a velocity gradient in the direction of migration and a plug-flow-like profile across the advancing sheet. The observed flow velocity can be decomposed into two contributions, a constant term stemming from directed cell

migration and a diffusion-like contribution that scales with the density gradient. From the density gradient and the speed of front propagation, we extract the collective diffusion coefficient of this diffusive component using the Fisher-Kolmogorov model. In order to connect diffusion mediated transport to underlying cellular motility, single cell trajectories and occurrence of vorticity were studied. We discovered that the directed large-scale cell flow alters fluctuations in cellular motion at short length scales: The formation of swirls is reduced compared to resting tissues. In addition, single-cell trajectories show persistent random-walk behavior superimposed on drift, whereas cells in resting tissue did not show significant displacement beyond neighboring cells. Our work thus suggests that active cell migration manifests itself in an underlying, spatially uniform drift as well as in randomized bursts of short-range correlated motion that lead to a diffusion-like transport.

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Lindlau	Jessica	
Ludwig	Stefan	LMU Munich
Mader	Andreas	LMU Munich
Manoharan	Muthiah	Alnylam Pharmaceuticals, Cambridge
Mast	Christof	LMU Munich
Milles	Lukas	LMU Munich
Milliron	Delia	University of Texas at Austin
Moghimi	Moein	University of Copenhagen
Morasch	Matthias	LMU Munich
Mühlbauer	Erika	LMU Munich
Müller	Jochen	LMU Munich
Nickel	Bert	LMU Munich
Novotny	Lukas	ETH Zürich
Palumbiny	Claudia Maria	TU Munich
Pill	Michael	Munich University of Applied Sciences
Pinto	Marilena	LMU Munich
Plank	Christian	TU Munich
Platonov	Sergey	LMU Munich
Rädler	Joachim	LMU Munich
Rehberg	Markus	LMU Munich
Reithmann	Emanuel	LMU Munich
Röder	Ruth	LMU Munich
Roller	Eva-Maria	LMU Munich
Saxena	Nitin	TU Munich
Schimmel	Dennis	LMU Munich
Schmidt	Alexandra	LMU Munich
Schollwöck	Ulrich	LMU Munich
Schreiber	Christoph	LMU Munich
Schuster	Simon	LMU Munich
Schwille	Petra	Max-Planck-Institute of Biochemistry
Sellier	Hermann	Institut NÈEL/CNRS. Grenoble
Sellner	Sabine	LMU Munich
Shi	Xian	LMU Munich
Stamp	Melanie	University of Augsburg
Stintzing	Sigmund	
Therv	Manuel	Honital Saint Louis Paris
Tinnefeld	Philin	
Urtel	Georg	
Utzinger	Maximilian	ethris GmbH
Verdorfer	Tobias	
von Dolft	lon	
Wagpor	Fract	
Wagner	Varanika	
Wengler	Daniala	
Werkmeister	Franz	
Westerbausen	Christoph	
Westermaise	Christian	
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Zeneumaier	Pene	
Znang	reng	
ZULU	matthias	

ACCOMMODATION

Accommodation for all participants is provided on San Servolo The buildings are situated in the island's beautiful green parkland. All bedrooms have air conditioning, television, telephone and internet access. A twenty-four hour reception service is guaranteed.

TIMETABLES

TRAIN TO VENICE AND BACK TO MUNICH

To Venic	e (21.09.)	Back to Munich (26.09.)						
Munich Main station	Venezia Santa Lucia	Venezia Santa Lucia	Munich Main station					
11:38	18:10	13:35	20:21					

LUNCH

The cafeteria is located on the ground floor of Building 15. The Cafeteria is open every day with the following timetable:

Breakfast	7.30 am - 9.30 am
Lunch	12.30 pm - 2.30 pm
Dinner	7.00 pm - 9.30 pm

Prices: breakfast € 6.50 (*please sign the list at the counter*), lunch and dinner € 10.00 (pasta course, main course, side order of vegetables or salad, yoghurt, bread and water). There are also reduced menus for € 7.00 (pasta course, side order of vegetables or salad, water and bread) and € 8.00 (main course, side order of vegetables or salad, water and bread).

A bar is also available on campus and is located on the ground floor of Area 6 in the main building.

WELCOME RECEPTION

The Welcome Reception will take place on Sunday, Sept. 21, at 8:00 pm.

INTERNET

WLAN Network San Servolo: UNIVIU

username: censworkshop2014 password: censworkshop2014

Two PC rooms with internet connection are accessible for the participants located next to the conference hall. Please ask for the keys in the conference office next to the lecture hall.

Internet activity will be monitored and recorded as required by Italian law.

BOAT LINE 20 TO WORKSHOP LOCATION (SAN SERVOLO) The boat from Venice to San Servolo leaves from the Riva degli Schiavoni at San Marco; the stop is in front of the Londra Palace Hotel. Boat number 20 goes to San Servolo.

To San	Servolo	Back to Venice						
S. Zaccaria	S. Servolo	S. Servolo	S. Zaccaria					
6:55	7:05	8:35	8:45					
7:15	7:25	8:45	8:55					
8:15	8:20	9:10	9:20					
8:35	8:50	9:40	9:50					
9:00	9:10	10:00	10:10					
9:20	9:30	10:50	11:00					
9:50	10:00	11:20	11:30					
10:30	10:40	12:10	12:20					
11:10	11:20	12:40	12:50					
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14:30	14:40	16:00	16:10					
15:10	15:20	16:50	17:00					
15:50	16:00	17:30	17:40					
		18:00	18:10					
every 40	min until	18:50	19:00					
		19:20	19:30					
20:30	20:40	20:10	20:20					
every h	our until	20:40	20:50					
23:30	23:40	21:50	22:00					
0:25	0:35	22:40	22:50					
1:30*	1:40							
2:10*	2:20							

* Trip made upon request. The boarding at S. Zaccaria must be booked at least 20 minutes beforehand at the following free phone number: (+39) 041-800-845065. Remember to arrive a few minutes before departure time.



	ime Friday, September 26	00:6	9:45 Florian Auras Covalent Organic Frameworks	0:05 Imaging electron transport in	semicontraction manustructures on the sectron wavelength	0:50 Closing remarks					Boat leaves at 11:20 / 12:10	Train to Munich leaves at	13:35 h from train station					
	Thursday, September 25 Ti	Moein Moghimi O Complement Sensing at Nanoscale	María García-Parajo Nanoscale imaging and single-mole- 00	tions using photonic antenna devices	COLLEE DIEAN	Liedewij Laan	0 How yeast adapts to a strong genetic perturbation: one function at the time	Alex Hamilton Strong spin-orbit coupling of spin-3/2 holes in gallium-arsenide semiconductor nanostructures	Lunch (12:30-14:00)	Charles Baroud	0 Microfluidic droplets for quantitative biological studies	5 Thomas Geislinger	Non-Inertial Lift Induced Cell Sorting		Posters session II & coffee	(00:71-60:61)	 Philip Tinnefeld Single-Molecule Fluorescence on DNA Origami 	Boat at 18:00
	Time	00:60	09:45	001	0.01		11:00	11:45			14:00	14:45					17:00	
	Wednesday, September 24	Stefan Ludwig 0.7-Anomaly in Quantum Point Contacts: Correlations in 1D	Hermann Sellier Electron Interactions in Quantum	Point Contacts		Dimitri Basov	Surface plasmons and phonon polaritons in atomically thin van der Waals crystals	Christian Plank Nucleic Acid Delivery – From Academic Discovery to Drug Development	Lunch (from 12:30) Boat at 12:40						Informal discussions			
	Time	00:60	09:45	00:01	00.01		11:00	11:45										
state values at the values of the values	Tuesday, September 23	Yaakov Benenson Molecular computing meets synthetic biology	Heinrich Leonhardt Studving cellular structure & function with	3D-SIM & fluorescent nanobodies		Alberto Amo	Quantum emulation with microcavity polaritons	Nynke Dekker Single-molecule studies of genome processing	Lunch (12:30-14:00)	Stephen Hart	Targeted nanocomplex formu- lations for gene and siRNA therapy	Christian Westermeier	Disorder in Org. Semiconductors		Posters session I &	сопее (15:05-17:00)	Lukas Novotny Cooling and Amplification of a Vacuum-Trapped Nanoparticle	18:00 Guided tour on San Servolo
nd Talk	Time	00:60	09:45	00.01	00.01		11:00	11:45			14:00	14:45					17:00	
Workshop Venice: Walk ar	Monday, September 22	Welcome Manuel Thery	Force ocaming in oness ribers Ralf Jungmann	Approaching the limit: Multiplexed Super- Resolution Microscopy with DNA-PAINT and Exchange-PAINT	Coffee break		Christof Mast Can thermal traps drive Darwinian evolution?	Phaedon Avouris Graphene Plasmons: Properties and Applications	Lunch (12:20-14:15)		Muthiah Manoharan Chemical Strategies for Delivery	of RNAi Drugs	Khaled Karrai	Precision positioning and sensing for nano-Manipulation applications	Coffee break	Delia Milliron Plasmonic metal oxide nanocrystals & their near infrared electrochromism	Peter Hänggi On the use and abuse of THERMODYNAMIC entropy	Boat at 18:00
CeNS	Time	09:00 09:15		10:00	10.45		11:15	11:35			14:15			15:00	15:45	16:15	17:00	